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Copyright (c) 1993 - 2000 Compugen Ltd.
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Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Human confluency r Human PRO1868 prot PRO1868, an A33 an Human PRO1868 protein e Secreted protein e Secreted protein e Human secreted pro	ID AAB27276 AAB33457 AAY96735 AAB80272 AAB80272 AAB80409 AAB80409 AAB80409 AAB38333 AAB38333 AAB383833 AAB383833 AAB383834 AAB80431	DB 211 211 221 222 222 222 222 222 222 22	Watch Length DB 100.0 310 2 100.0 310 2 100.0 310 2 100.0 310 2 100.0 310 2 100.0 310 2 100.0 310 2 100.0 310 2 100.0 310 2 100.0 311 2 100.0 311 2	Match 100.0 100.0 100.0 100.0 100.0 100.0 100.0 100.0	Score 1637 1637 1637 1637 1637 1637 1637 1637	Result NO: 1 2 3 3 4 4 5 5 6 6 7 7 8 9 9 9
Gene #13 associate	AAB80431	22	339	100.0	1637	11
Human secreted pro	AAB38384	21	311	100.0	1637	10
Human secreted pro	AAB38383	21	311	100.0	1637	و
Human secreted pro	AAB38333	21	311	100.0	1637	8
Secreted protein e	AAB80409	22	310	100.0	1637	7
Secreted protein e	AAB80408	22	310	100.0	1637	6
Secreted protein e	AAB80383	22	310	100.0	1637	u
Human PRO1868 prot	AAB80272	22	310	100.0	1637	4
PRO1868, an A33 an	AAY96735	21	310	100.0	1637	ω
Human PRO1868 prot	AAB33457	21	310	100.0	1637	2
Human confluency r	AAB27276	21	310	100.0	1637	-
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AAB55950	AAB80232	AAB31202	AAY76076	AAY76011	AAY70670	AAY95344	AAB24405	AAW74464	AAY13364	AAY23321	AAY08071	AAB39253	AAB50904	AAY23325	AAW61380	AAY11472	AAB53081	AAB80222	AAY70668	AAB24401	AAB33421	AAY13354	AAY23324	AAY08060	AAB27275	AAB27273	AAW75220	AAU00512	AAW85457	AAB27278	AAB27272	AAB39254	AAY96294
Skin cell protein,	Human PRO301 prote	Amino acid sequenc	Human A33 receptor	Human A33 receptor	Human PRO301 prote		Human PRO301 prote	F11 antigen protei	Amino acid sequenc	Amino acid sequenc	Human PRO307 prote	Gene 15 human secr	Human PRO245 prote	 A33 related antige 	Mouse junctional a	Human 5' EST secre	gioge	Human PRO245 prote			Human PRO245 prote	Amino acid sequenc	A33 related antige	Human PRO245 prote	Murine confluency	Human confluency r	Human secreted pro	Human junctional a	Secreted protein e	Murine confluency	Human confluency r	Human secreted pro	Human IGFAM-6 immu

ALIGNMENTS

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CRAM-2) polypeptide, useful for treatment of tumors, inflammation reactions and modulating vascular permeability -	Isolated human Confluency Regulated Adhesion Molecule 1 or 2 (CRAM-1		N-PSDB; AAA95306.	WPI; 2000-587436/55.		Imhof BA, Aurrand-Lions M;		(RMFD-) RMF DICTAGENE SA.		11-MAR-1999; 99EP-0200746.		13-MAR-2000; 2000WO-EP02219.		14-SEP-2000.		WO200053749-A2.		Homo sapiens.		confluency regulated adhesion molecule 1; CRAM-1; JAM-2.	inflammation; cancer; wound; angiogenesis; human;	Immunoqlobulin superfamily; Iq Sf; vascular adhesion molecule;		Human confluency regulated adhesion molecule 1 #2.		23-FEB-2001 (first entry)		AAB27276;		AAB27276 standard; Protein; 310 AA.	RESULT 1

Isolated human Confluency Regulated Adhesion Molecule 1 or 2 (CRAM-1 or CRAM-2) polypeptide, useful for treatment of tumors, inflammation reactions and modulating vascular permeability -

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                                                                                                              dermatological; antiarthritic; antirheumatic; immunosuppressive; hamoustatic; antithyroid; antidiabetic; noctropic; neuroprotective; antianaemic; hepatotropic; virucide; antipsoriatic; antiallergic; antiasthmatic; systemic lupus erythematosus; rheumatoid arthritis; osteoarthritis; spondyloarthropathy; systemic sclerosis; sarcoidosis; idiopathic inflammatory myopathy; Sjogren's syndrome; thyroiditis; systemic vasculitis; autoimmune haemolytic anaemia; diabetes mellitus; autoimmune thrombocytopaenia; immune-mediated renal disease; demyelinating disease; hepatobiliary disease; Whipple's disease; inflammatory bowel disease; glutten-sensitive enteropathy; autoimmune disease; immune-mediated skin disease; allergic disease; immunological disease; transplantation associated disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence is the human confluency regulated adhesion molecule I (CRAM-I, also known as JAM-2). CRAM-I is one of the vascular adhesion proteins of the immunoglobulin superfamily (Ig Sf). The CRAM-I protein and coding sequence can be used in the treatment of cancer, inflammation, to modulate cell-cell interactions and angiogenesis, and in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2;
    14-SEP 2000.
                                                                                                                                                                                                                                                                                                                                                                                                AAB33457;
                                                                                                                                                                                                                                                                                                                                                                                                                            AAB33457 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modulation of wound healing.
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                                                                        sapiens.
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                                                                                                     rejection;
                                                                                                                                                                                                                                                                                                      immune related disease; diagnosis; antiinflammatory; cardiant;
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                                                                                                     graft-versus-host-disease
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The present invention describes sixty four human PRO proteins which can CC be used in the treatment of immune related diseases. The human PRO CC proteins, anti-PRO antibodies, agonists and antagonists are useful for CC treating and diagnosing immune related disorders. The disorders are CC selected from systemic lupus erythematosus, rheumatoid arthritis, constearthritis, juvenile chronic arthritis, spondyloarthropathies, Systemic sclerosis, idiopathic inflammatory myopathies, Sjogren's CC syndrome, systemic vasculitis, sarcoidosis, autoimmune haemolytic anaemia, autoimmune thrombocytopaenia, thyroiditis, diabetes mellitus, CC immune-mediated renal disease, demyelinating diseases of the central CC and peripheral nervous systems, hepatobiliary diseases, inflammatory CC bowel disease, gluten-sensitive enteropathy and Whipple's disease, CC autoimmune or immune-mediated skin diseases, allergic diseases, CC diseases including graft rejection and graft-versus-host-disease.
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20-DEC-1999;
30-DEC-1999;
30-DEC-1999;
05-JAN-2000;
06-JAN-2000;
06-JAN-2000;
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05-OCT-1999;
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29-NOV-1999;
30-NOV-1999;
31-NOV-1999;
01-DEC-1999;
01-DEC-1999;
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01-SEP-1999;
08-SEP-1999;
13-SEP-1999;
                                                                                                                                                                                                                                          Sixty four PRO polypeptides, useful in the diagnosis and treatment immune related disorders, e.g. systemic lupus erythematosis, rheuma arthritis, osteoarthritis, thyroiditis and diabetes mellitus - \frac{1}{2}
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02-JUN-1999
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N-PSDB; AAC58622.
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Kabakoff RC,
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Best Local
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                                                                                                                                                                                                                                                           PRO1868; A33 antigen; secreted protein; transmembrane protein; anti-inflammatory; cytostatic; recombinant production; gene the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         in the isolation of human PRO sequences. AAC58579 to AAC58642 and AAB33414 to AAB33477 represent human PRO polynucleotide and protein sequences given in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    DFRHKSSFVI 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PTDSRANPRFRNSSFHLNSETGTLVFTAVHKDDSGQYYCIASNDAGSARCEEQEMEVYDL
                                                                                                                                                                                                                                                                                                                                                                                                               dfrhkssfvi 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ptdsranprfrnssfhlnsetgtlvftavhkddsggyyciasndagsarceeqemevydl
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                                                                                                                       /note- "N
106..109
                            /note-
226..2
                                                                                                    /note-
107..1
                                                                                                                                                                          26..31
/note= "N-myristoylation site"
                                                                      192..195
                                                                                                                                             /note= "Tyrosine kinase phosphorylation site"
104..107
                                                                                                                                                                                               /label= Signal_peptide
                                                                                                                                                                                                                     Location/Qualifiers
                             . 231
                                                          "N-glycosylation
                                                                                           CAMP-
                                                                                                           *Casein kinase II phosphorylation
                                                                                                                                  "N-glycosylation
                  "N-myristoylation site"
                                      "N-myristoylation site"
                                                                               'cAMP- and cGMP-dependent protein kinase
phosphorylation site"
                                                                                                                                                                                                                                                                                           homologue
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Pred. No. 1
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                                                           site"
                                                                                                                                  site"
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l.2e-133;
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Best Local Similarity
Matches 310; Conserv
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12-FEB-1999;
02-JUN-1999;
                                                                                                                                                                                                                                                                 This is PRO1868, a putative homologue of A33 antigen, a known colorectal cancer-associated marker. The invention concerns novel secreted and transmembrane proteins, designated PRO polypeptides. The cDNA and gene sequences are useful in the recombinant production of PRO polypeptides, as a hybridization probe to screen libraries to isolate cDNAs with sequence identity to PRO polypeptides or to map the gene encoding the PRO polypeptides and analyzing genetic disorders. The cDNA/gene can also be used to produce transgenic animals useful for the development and screening of therapeutically useful reagents. They can also be used in gene therapy, e.g. to replace a defective gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-431586/37.
N-PSDB; AAA51265.
                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                       transmembrane polypeptide
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12-JAN-1999;
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Ood
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12-JAN-1999;
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16-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated nucleic acid molecule encodes a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GETH ) GENENTECH INC.
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                       181
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 181
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                                                                                                                                      SDPRIEWKKIQDEQTTYVFFDNKIQGDLAGRAEILGKTSLKIWNVTRRDSALYRCEVVAR 120
ptdsranprfrnssfhlnsetgtlvftavhkddsggyyciasndagsarceeqemevydl
          PTDSRANPRFRNSSFHLNSETGTLVFTAVHKDDSGQYYCIASNDAGSARCEEQEMEVYDL
                                           ndrkeideivieltvqvkpvtpvcrvpkavpvgkmatlhcqeseghprphyswyrndvpl
                                                         NDRKEIDEIVIELTVQVKPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYSWYRNDVPL
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                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                14; 154pp;
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an J, Roy MA,
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9805-0115145

9805-011551

9805-011558

9905-011558

9905-011573

9905-011573

9905-0119341

9905-0119345

9905-0119365

9900-0812252
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/note= "N
244..249
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                                                                                                                                                                                                100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                English.
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                                                                                                                                                                                    0
                                                                                                                                                                                               Score 1637; DB 21; Pred. No. 1.2e-133;
                                                                                                                                                                                    Mismatches
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D, Watanabe CK;
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                                                                                                                                                                                                           310;
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                                                                                                                                                                        Shkenazi AJ, Bouse
Filvaroff E, Fong S
Godowski PJ, Grima)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Auman; PRO; dermatological; anti
antiparkinsonian nootropic; neur
antiangiogenic; vasotropic; anti-
ntiarthritic; antiinfertility;
ophthalmological; gene therapy;
                                                                                          Sixty one nucleic acids encoding PRO polypeptides which are useful in the treatment of skin diseases (e.g. psoriasis), cancers (e.g. lung squamous cell carcinoma) and neurodegenerative diseases (e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo
enterocolitis),
          The present sequence is one of sixty one novel secreted and transmembrane PRO polypeptides. The PRO polypeptides are useful for treating skin diseases (e.g. psoriasis), cancers (e.g. squamous cell carcinoma), gastrointestinal disorders (e.g.
                                                              Claim 1; Fig
                                                                                  Alzheimer's disease)
                                                                                                                                      N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                 26-JUL-1999;
28-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                     07-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ischaemia; inflammation.
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                                                              124;
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                                                                                                                                                                            Fong S, Gao W, Gert
Grimaldi CJ, Gurney
an J, Paoni NF, Roy
                                                                                                                                                                                                                                                                                                                                                                                                           2000WO-US04414
                                                                                                                                                                   Wood WI;
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99WO-US30999.
99WO-US00219.
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99WO-US23089.
99WO-US28214.
99WO-US28313.
                                                                                                                                                                                                           Botstein D,
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                                                             393pp;
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                                                              English.
                                                                                                                                                                            Desnoyers L,
W, Gerber H,
Gurney AL, H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antipsoriatic; cytostatic; antiinflammatory; neuroprotective; vulnerary; cardiant; antiasthmatic; antirheumatic; cancer; .ty; antidiabetic; antiviral; diabetes; apy; skin disease; gastrointestinal disorder;
 diseases
                                                                                                                                                                            Hillan KJ,
Stewart TA,
                                                                                                                                                                                                          Eaton DL,
(e.g. Alzheimer's
         disorders (e.g
                                                                                                                                                                             Kljavin IJ;
Tumas D;
                                                                                                                                                                                                          Ferrara N;
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Best Local
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diagnosing, preventing hyperproliferative, can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Secreted protein; cardiovascular; c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Secreted
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                         New isolated nucleic acid encoding 29 secreted proteins, diagnosing, preventing and treating e.g. autoimmune,
                                                                                                   WPI; 2001-123261/13
                                                                                                                                                                                                                         (HUMA-) HUMAN GENOME
                                                                                                                                                                                                                                                                                                                          20-JUL-2000;
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1 dfrhkssfvi
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                                                                                                                                                 CA, Ruben
, Lafleur
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein encoded by
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                                                                                                                                                                                                                                                                            99US-0145220
                                                                                                                                                 n SM,
DW,
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  cardiovascular,
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Pred. No. 1.2e-133;
; Mismatches 0;
                                                                                                                                                 Duan RD,
Birse CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ine; hyperproliferation;
infection; food.
  e.g. autoimmune, and ocular disc.
                                                                                                                                                 Ni J, Soppet DR,
Komatsoulis GA;
  diseases or disorders
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                                                                                                                                                                                                                                        AAB80408 standard; protein; 310
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                                                                                                                                                    Secreted protein; human; autoimmune; hyperproliferation; cardiovascular; cerebrovascular; infection; food.
                                    (HUMA-) HUMAN GENOME SCI INC
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  n CA, ....
v. Lafleur
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         Ruben SM,
afleur DW,
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Komatsoulis GA;
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                        23-JUL-1999;
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                                                            Immunosuppressive; antiarthritic; ar cytostatic; cardiant; vasotropic; ce nootropic; antibacterial; virucide;
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                                                vulnerary; gene
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HS,
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Pred. No. 1.2e-133;
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Birse CE,
                                                            cerebroprotective; neuroprotec
e; fungicide; opthalmalogical;
                                                                                                                  gene 13 clone HAPSA79
                                                                                         antirheumatic;
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Komatsoulis GA;
                                                                                       antiproliferative;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC Sequences AAB38321-B38396 represent the amino acid sequences of 62 CC human secreted proteins encoded by the genes AAC69512-C69587. The genes CC and proteins are useful for preventing, ameliorating or treating medical CC conditions, e.g. by protein or gene therapy. The genes are isolated from CC a range of human tissues disclosed in the specification. The nucleic CC claignosis, antibodies and (ant)agonists are useful in the CC diagnosis, treatment and prevention of: (a) autoimmune diseases e.g. CC rheumatoid arthritis; (b) hyperproliferative disorders e.g. neoplasms CC of the breast or liver; (c) cardiovascular disorders e.g. cardiac CC arrest; (d) cerebrovascular disorders e.g. cerebral ischemia; (e) CC anglogenesis; (f) nervous system disorders e.g. cerebral ischemia; (e) anglogenesis; (f) nervous system disorders e.g. Alzheimer's disease; (g) cinfections caused by bacteria, viruses and fungi; and (h) ocular CC disorders e.g. corneal infection. The polypeptides can also be used to CC aid wound healing and epithelial cell proliferation, to prevent skin CC supporting cell culture of primary tissues, to regenerate tissues and in
                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local Similarity
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26-APR-1999;
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                              DFRHKSSFVI 310
                                                                              NIGGIIGGVLVVLAVLALITLGICCAYRRGYFINNKQDGESYKNPGKPDGVNYIRTDEEG
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                                                             niggiiggvlvvlavlalitlgiccayrrgyfinnkqdgesyknpgkpdgvnyirtdeeg
                                                                                                                              ptdsranprfrnssfhlnsetgtlvftavhkddsgqyyciasndagsarceeqemevydl
                                                                                                                                             PTDSRANPRFRNSSFHLNSETGTLVFTAVHKDDSGQYYCIASNDAGSARCEEQEMEVYDL
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99US-0130991.
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n HS, Ebner
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Best Local
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Lafleur DW,
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26-APR-1999;
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Ni Olsen
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Pred. No. 1.2e-133;
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                      Sequences AAB38321-B38396 represent the amino acid sequences of 62 CC human secreted proteins encoded by the genes AAC69512-C69587. The genes CC and proteins are useful for preventing, ameliorating or treating medical conditions, e.g. by protein or gene therapy. The genes are isolated from CC arios, proteins, antibodies and (ant)agonists are useful in the CC diagnosis, treatment and prevention of: (a) autoimune diseases e.g. creating the present or liver; (b) hyperproliferative disorders e.g. neoplasms CC of the breast or liver; (c) cardiovascular disorders e.g. cardiac carrest; (d) cerebrovascular disorders e.g. cerebral isohemia; (e) cardiovascular disorders e.g. cardiac carest; (d) cerebrovascular disorders e.g. Alzheimer's disease; (g) cinfections caused by bacteria, viruses and fungi; and (h) ocular CC disorders e.g. corneal infection. The polypeptides can also be used to augung due to sunburn, to maintain organs before transplantation, for carearers entraced.
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                                                                                                                                                                                                                                                                                                                                                                                                            Ruben SM, N
Lafleur DW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Immunosuppressive; antiarthritic; antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; neuroprotectiv nootropic; antibacterial; virucide; fungicide; opthalmalogical; hum
supporting chemotaxis
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26-APR-1999;
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99US-0130991.
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n HS, Ebner
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                                                                                                                                                                                                                                                                                                                                                                                                               Rosen CA, S
Florence KA,
               to regenerate tissues
                                                                                                                                                                                                                                                                                                                                                                                                               Soppet DR,
, Moore PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                               Shi Y;
Birse
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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
invention is used to prevent autoimmune diseases e.g. arthritis, hyperproliferative disorders e.g. neoplasms
                                                                                                     Disclosure; Page 75; 601pp; English.
                                                                                                                                                      New isolated nucleic acid encoding 29 secreted proteins, diagnosing, preventing and treating e.g. autoimmune, hyperproliferative, cardiovascular, and ocular diseases and microorganism infections
                                                                                                                                                                                                                                                                                                                                       Rosen CA, Ruben SM,
Shi Y, Lafleur DW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Secreted protein; cardiovascular; c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gene #13 associated peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                     (HUMA-) HUMAN GENOME SCI INC
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|||||||||
1 dfrhkssfvi 310
                             present invention relates to 29 human secreted proteins. The ention is used to prevent autoimmune diseases e.g. rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NDRKEIDEIVIELTVQVKPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYSWYRNDVPL 180
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                                                                                                                                                                                                                                                                                                                                          Ebner R,
Olsen HS,
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                                                                                                                                                                                                                                                                                                                                          Duan RD,
Birse CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hyperproliferation;
                                                                                                                                                                                   ocular diseases or disorders
                                                                                                                                                                                                                                                                                                                                          Ni J, Soppet DR, Komatsoulis GA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                                                                                                                                                                                                                                    Moore PA;
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                                                                                                              RESULT 12
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischemia, angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. Also used in food preparations.
                                                                                                                                                                                               Key
                                                                                                                                                                                                                                                   Human;
                                                                                                                                                                                                                                                                      Human IGFAM-6 immunoglobulin
19-NOV-1998;
22-DEC-1998;
                                                                                                      Domain
                                                                                                                         Domain
                                                                                                                                             Domain
                                                                                                                                                                 Protein
                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                        infection;
                                                                                                                                                                                                                                                                                           16-AUG-2000
                                                                                                                                                                                                                                                                                                                                   AAY96294 standard; protein; 310
                               19-NOV-1999;
                                                   25-MAY-2000
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                                                                                                                                                                                                                                        immunoglobulin; IGFAM-6; IGFAM; immune disorder;
ion; inflammation; haematopoiesis; AIDS; allergy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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98US-0113635
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                                                                                                      /label=
238..260
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                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                      /label=
                                                                                                                                    label-
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                                                                                                             Ig_domain
                                                                                                                                   Ig_domain
                                                                                            transmembrane_domain
                                                                                                                                                       IGFAM-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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                                                                                                                                                                                                                                                   cancer;
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RESULT 1
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Best Local.S
Matches 309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tissue. It is expressed in reproductive, nervous and reardiovascular tissue, where cancer and inflammation are common. The yene, protein, its antibodies, agonists and antagonists are suitable for diagnosing and treating many diseases, including cancer, immune system disorders (such as inflammation, AIDS, allergies, anaemia, arteriosclerosis, asthma, atherosclerosis, cholecystitis, Crohn's disease, diabetes mellitus, emphysema, Graves' disease, hepatitis, multiple sclerosis, psoriasis, rheumatoid arthritis, scleroderma, systemic lupus erythematosus and ulcerative colitis, complications of cancer, haemodialysis and extracorporeal circulation, trauma and haematopoietic cancer (such as leukaemia) and infections caused by bacteria, viruses, fungi or parasites.
                         02-FEB-2001
                                                                      AAB39254 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunoglobulin superfamily proteins, i protein is useful for preventing and altered levels of the protein such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ę
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                                                                                                                                  301
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                                                                                                                                              DFRHKSSFVI 310
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                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                         (first entry)
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protein sequence encoded
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Hillman JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                      99.58;
                                                                                                                                                                                                                                                                                                                                                                                                                            <u>..</u>
                                                                                                                                                                                                                                                                                                                                                                                                                           Score 1629; DB 21;
Pred. No. 6.1e-133;
0; Mismatches 1;
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Yang J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the agonist and antagonist of treating disorders associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cancer,
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by gene
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15 SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 310;
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GCLIGAVNLKSSNRTPVVQEFESVELSCIITDSQTSDPRIEWKKIQDEQTTYVFFDNKIQ 85

gcligavnlkssnrtpvvqefesvelsciitdsqtsdpriewkkiqdeqttyvffdnkiq

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GDLAGRAEILGKTSLKIWNVTRRDSALYRCEVVARNDRKEIDEIVIELTVQVKPVTPVCR 145 gdlagraeilgktslkiwnvtrrdsalyrcevvarndrkxideivieltvqvkpvtpvcr 120

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121 146

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The polynucleotide sequences given in AAC74223-C74279 encode the human consists and proteins represented in AAB39179-B3926. Sequences and also protein sequences with which they share homology. The proteins have activities based on the tissues and cells in which they are expressed. Examples of activities include: immunosuppressive; antiarthritic; antiproliferative; cytostatic; cardiant; vasotropic; cc antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cc polynucleotide; and opthalmalogical. The human secreted proteins, cc polynucleotides, antagonists and agonists of the invention may be useful in the treatment, prevention, and/or diagnosis of various disease, disorders and conditions such as autoimmune diseases e.g. rheumatoid cc arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cardiac arrest, cerebrovascular cc e.g. Alzheimer's disease, infections caused by bacteria, viruses and cc e.g. Alzheimer's disease, maintain organs before transplantation, in cc regenerate tissues, maintain organs before transplantation, in cc chemotaxis and as a food additive or preservative e.g. to horcease con during the isolation and characterisation of the genes of the invention.
Query Match
Best Local S
Matches 283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; secreted protein; immunosuppressive; antiarthritic; antirheumatic antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; neoplasm opthalmalogical; autoimmune disease; rheumatoid arthritis; angiogenesis; hyperproliferative disorder; cardiovascular disorder; infection; cerebrovascular disorder; nervous system disorder; ocular disorder; wound healing; chemotaxis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Isolated nucleic acid molecule encoding a human secreted protein used in preventing, treating or ameliorating a medical condition
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10-DEC-1999;
                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-MAR-2000; 2000WO-US06792
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (HUMA-) HUMAN GENOME
                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAC74237
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 32;
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99US-0169980.
                                   91.5%;
99.3%;
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Score 1498; D
Pred. No. 1.1e
0; Mismatches
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; DB 21;
1.1e-121;
hes 2;
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                                                                                                                                            Matches 266;
                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                   The
                                                                                                                                                                                                                       (CRAM-1, also known as JAM-2). CRAM-1 is one of the vascular adhesion roteins of the immunoglobulin superfamily (Ig Sf). The CRAM-1 protein and coding sequence can be used in the treatment of cancer, inflammation, to modulate cell-cell interactions and angiogenesis, and
                                                                                                                                                                                                                                                                                    Claim 1; Fig 3; 59pp; English.
                                                                                                                                                                                                                                                                                                      Isolated human Confluency Regulated Adhesion Molecule 1 or 2 (CRAM-1 or CRAM-2) polypeptide, useful for treatment of tumors, inflammation reactions and modulating vascular permeability -
                                                                                                                                                                                                                                                                                                                                                                                                            11-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                13-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mmunoglobulin superfamily; Ig Sf; vascular adhesion molecule; inflammation; cancer; wound; angiogenesis; human; confluency regulated adhesion molecule 1; CRAM-1; JAM-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB27272 standard; Protein; 310
                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                     Imhof BA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human confluency regulated adhesion molecule 1 #1.
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                     121
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                                                                                                                                           Local Similarity 85.
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                                                                                                  the modulation of
                                                                                                                                                                                                                                                                 present sequence is the human confluency regulated adhesion molecule
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                                                                                                                                                                                                                                                                                                                                                 2000-587436/55
 PTDSRANPRFRNSSFHLNSETGTLVFTAVHKDDSGQYYCIASNDAGSARCEEQEMEVYDL
                   Aurrand-Lions M;
                                                                                                                                                                                             310 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                           99EP-0200746
                                                                                                                                                     86.1%;
85.8%;
                                                                                                                                                                                                                wound healing.
                                                                                                                                           18;
                                                                                                                                          Score 1409; DB 21;
Pred. No. 6.3e-114;
8; Mismatches 26;
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                                                                                                                                                             Length
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RESULT 1
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                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                       The present sequence is the murine confluency regulated adhesion molecule 1 (CRAM-1, also known as JAM-2). CRAM-1 is one of the vascular adhesion proteins of the immunoglobulin superfamily (Ig Sf). The CRAM-1 protein and coding sequence can be used in the treatment of cancer, inflammation, to modulate cell-cell interactions and angiogenesis, and in the modulation of wound healing.
                                                                                                                                                                                                                                                                                     Isolated human Confluency Regulated Adhesion Molecule 1 or 2 (CRAM-1 CRAM-2) polypeptide, useful for treatment of tumors, inflammation reactions and modulating vascular permeability -
                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                N-PSDB; AAA97189.
                                                                                                                                                                                                                                                                                                                                                                                                                              13-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               confluency regulated adhesion molecule 1; CRAM-1; JAM-2
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                                                                                                                                                                                                                                                                   Example; Fig 8; 59pp; English.
                                                                                                                                                                                                                                                                                                                                           WPI; 2000-587436/55
                                                                                                                                                                                                                                                                                                                                                                Imhof BA,
                                                                                                                                                                                                                                                                                                                                                                                                          11-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inflammation; cancer; wound; angiogenesis; mouse;
                                                                                                                                                                                                                                                                                                                                                                                    (RMFD-) RMF DICTAGENE SA.
           121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mmunoglobulin superfamily;
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NDRKEIDEIVIELTVQVKPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYSWYRNDVPL
                             SDPRIEWKKIQDEQTTYVFFDNKIQGDLAGRAEILGKTSLKIWNVTRRDSALYRCEVVAR 120
                                                                      NIGGIIGGVLVVLAVLALITLGICCAYRRGYFINNKQDGESYKNPGKPDGVNYIRTDEEG 300
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                                                                                                                   Conservative
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                                                                                                                            Score 1409; DB 21;
Pred. No. 6.3e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sf; vascular adhesion molecule;
                                                                                                                   Mismatches
                                                                                                                                     Length 310;
                                                                                                                   Indels
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                                                                                                                                                                                   human foetal pineal gland
(e.g. as vaccines) or suppressing activity, haematopoiesis regulating activity, tissue growth activity, activin/inhibin activity, chemotactic/chemokinetic activity, haematostatic and thrombolytic activity, receptor/ligand activity, anti-inflammatory activity, cadherin/tumour invasion suppressor activity, and tumour inhibition activity (no data is given in the specification to support these activities). The
                                                                              The present sequence represents a secreted protein. The polynucleotide and secreted protein are predicted to have biological activities which would make them Suitable for treating, preventing or ameliorating medical conditions in humans and animals, although no supporting data is given. Suggested activities include nutritional activity, immune stimulating
                                                                                                                                                                                       New polynucleotides encoding secreted human proteins - derived from human foetal brain, adult brain, foetal kidney, placenta or adult pineal gland cDNA libraries.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 suppressing activity; haematopolesis regulating activity; tissue growth activity; activin; inhibin activity; chemotactaxis; chemokinetic activity; haemostasis; thrombolytic activity; receptor; ligand; anti-inflammatory; cadherin; tumour invasion suppressor;
                                                                                                                                                              Claim 17; Page 73-74; 113pp; English
                                                                                                                                                                                                                                            WPI; 1998-609890/51.
N-PSDB; AAV82780.
                                                                                                                                                                                                                                                                                                                                                       19-MAR-1998;
21-MAR-1997;
                                                                                                                                                                                                                                                                                     Racie
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tumour inhibition;
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Spaulding
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97US-0822167.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nutritional activity; immune stimulating;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Protein;
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RESULT 17
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Pred. No. 2.4e-34;
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                           diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaen developmental abnormality; foetal deficiency; blood; allergy; renal; immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma; inflammation; ischaemic shock; Alzheimer's disease; restenosis; AIDS;
osteoporosis;
             cognitive disorder; schizophrenia; prostate; obesity;
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for producing antibodies that are suitable for therapeutic
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                                                                                              secreted protein; fusion protein; gene therapy; protein therapy
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Best Local
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                                                                                                                                                                                                                                                                                                                                This sequence represents a secreted human protein encoded by the gene clone detailed in the descriptor line. The gene can be used to generate fusion proteins by linking to the gene to a human immunoglobulin Fc portion (e.g. AAV34277) for increasing the stability of the fused protein as compared to the human protein only. The invention relates to 28 novel genes and their fragments (nucleic acid sequences: AAV34286-V34325; amino acid sequences AAW75196-W75235) which are useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. Also, pathological conditions can be diagnosed by determining the amount of the new polynucleotides. Specific uses are described for each of the 28 polynucleotides, based on which tissues they are most highly expressed in the new polynucleotides. Specific uses are described for each of the 28 polynucleotides, based on which tissues they are most highly expressed in the account of the case o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ferrie AM,
Li H, Li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-PSDB;
                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 168-169; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isolated human poly:nucleotide(s) encoding secretory peptide(s) used to develop products for the diagnosis and treatment of e.g inflammation, cancers, CNS disorders or immune system disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-SEP-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             endocrine; metabolism; regulation; malabsorption;
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                                                                                                                                                                                                                                                                                                                     see AAV34286
   114
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                                                                                                                                                                                                       Local Similarity
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RCEVVARNDR-KEIDEIVIELTVQVKPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1998-520811/44
DB; AAV34310.
                                                                                                                                                   MALRRPPRIRICARLPDFFLLLLFRGCLIG-----AVNLKSSNRTPVVQ--EFESVELSC
                                    -ktpkktvxsrlewkkl-grsvsfvyyqqtlqqdfknraemi-dfniriknvtrsdagky
                                                                          IITDSQTSDPRIEWKKIQDEQTTYVFFDNKIQGDLAGRAEILGKTSLKIWNVTRRDSALY
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                                                                                                                                                                                                                                                                                298 AA;
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                                                                                                                                                                                        Conservative
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97US-0048357.
97US-0050934.
97US-0048970.
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97US-0040710.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /label= unknown
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                                                                                                                                                                                                                                                                                                                   described uses) .:
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                                                                                                                                                                                      60;
                                                                                                                                                                                      Score 481; DB 19;
Pred. No. 9.7e-34;
0; Mismatches .116
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Ruben SM,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    , Kyaw H;
Soppet DR;
                                                                                                                                                                                                                          Length 298;
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ulin Fc
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                                                                                                                                          RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
21-JUN-1999
                                             AAY11472;
                                                                                          AAY11472 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              to map genes encoding PRO proteins, to analyse genetic disorders, and gene therapy. PRO nucleic acids can also be used to produce transgenic animals useful for the development and screening of potential therapeutic agents. The present sequence represents a PRO protein of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      comprising
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 designated PRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 69; Fig 36; 293pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       angiogenic disorders, such as atherosclerosis, wounds or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New isolated nucleic acid for producing a PRO polypeptide, analyzing genetic disorders and treating cardiovascular, endothelial or
                                                                                                                                                                                                                                                                                                                168
                                                                                                                                                                                                                                                                                                                                                                173
                                                                                                                                                                                                                                                                                                                                                                                                               108
                                                                                                                                                                                                                                                                                                                                                                                                                                                          114
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 marrsrhrl-------lllllrylvvalgyhkaygfsapkdqqvvtaveyqeailac 50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   proteins. The invention also relates to vectors and host cells prising a PRO nucleic acid, the recombinant production of a PRO tein, PRO antibodies specific for a PRO protein, fusion proteins
                                                                                                                                                                                                              krmqvddlnisgiiaavvvvalvisvcglgvcyaqrkgyf 267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IITDSQTSDPRIEWKKIQDEQTTYVFFDNKIQGDLAGRAEILGKTSLKIWNVTRRDSALY 113
                                                                                                                                                                                                                                                                                                                                                                                                                                   RCEVVARNDR-KEIDEIVIELTVQVKPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYS 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -ktpkktvssrlewkkl-grsvsfvyyqqtlqgdfknraemi-dfniriknvtrsdagky 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         104;
                                                                                                                                                                                                                                                                                                                                                           WYRNDVPLPTDSRANPRFRNSSFHLNSETGTLVFTAVHKDDSGQYYCIASNDAGSARCEE 232
                                                                                                                                                                                                                                                             QEMEVYDLNIGGIIGGVLVVLAVLALITLGICCAYRRGYF 272
                                                                                                                                                                                                                                                                                                        {\tt wfkdgirllenprlgsqstnssytmntktgtlqfntvskldtgeyscearnsvgyrrcpg}
                                                                                                                                                                                                                                                                                                                                                                                                          rcevsapseggqnleedtvtlevlvapavpscevpssalsgtvvelrcqdkegnpapeyt
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28.2%; Score 461.5; DB 22; 37.1%; Pred. No. 5e-32;
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RESULT :
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       upstream regulatory sequence; cytokine activity; cell proliferation; differentiation; haematopolesis regulation; tissue growth regulation; treproductive hormone regulation; chemotactic; chemokination; thrombolytic; anti-inflammata-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human secreted proteins, and encode the proteins given in AAV11374 to AAV11371, respectively. The proteins given represent the signal peptide and an N-terminal fragment of a secreted protein. The nucleic acid sequences can be used for producing secreted human gene products. They can also be used to develop products for diagnosis and therapy. The
AAW61380
                          AAW61380 standard;
                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX39440 to AAX39597 represent 5' expressed sequence tags (ESTs) for human secreted proteins, and encode the proteins given in AAX11374 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 34; Page 394; 434pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and fetal brain tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New nucleic acids encoding human cDNA libraries prepared from subs
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                                                                                                    61 sdpriewkkiqdeqttyvffdnkiqgdla
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                                                                                                                                                                                                                       28.1%; ilarity 100.0%; Conservative
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                         protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        English.
                                                                                                                                                                                                                       Score 460; Db zv;
; Pred. No. 1.3e-32;
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Matches 104; Conserv
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Godowski PJ, Gri
                                                                                                                                                                The present sequence is one of sixty one novel secreted and transmembrane PRO polypeptides. The PRO polypeptides are useful for treating skin diseases (e.g. psoriasis), cancers (e.g. lung squamous cell carcinoma), gastrointestinal disorders (e.g. enterocolitis), neurodegenerative diseases (e.g. Alzheimer's disease, parkinson's disease), wound repair, cardiovascular disorders (e.g. parkinson's disease), wound repair, cardiovascular disorders (e.g. endometrial bleeding angiogenesis, ischaemias such as coronary ischaemia, atherosclerosis), inflammatory disorders (e.g. asthma, rheumatoid arthritis, multiple sclerosis), infertility, AIDS and diabetes and retinal disorders such as retinitis pigmentosum. The PRO nucleic acids have applications in molecular biology, including use as hybridization probes, and in chromosome and gene mapping.
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20-DEC-1999;
20-DEC-1999;
05-JAN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sixty one nucleic acids encoding PRO polypeptides which are useful in the treatment of skin diseases (e.g. psoriasis), cancers (e.g. lung lamous cell carcinoma) and neurodegenerative diseases (e.g. theimer's disease)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ashkenazi AJ,
Filvaroff E,
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N-PSDB; AAF72383.
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05-OCT-1999;
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08-SEP-1999;
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Fong S, Gao W, Gerber H,
Grimaldi CJ, Gurney AL, F
an J, Paoni NF, Roy MA, S
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99US-0146222.
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99WO-US00219.
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                            28.2%;
     52;
                         Score 461.5; DB
Pred. No. 5e-32;
       Mismatches
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Hillan KJ,
Stewart TA,
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                                                   DB 22;
       103;
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Tumas D;
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ME, Goddai
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       21;
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01-SEP-1999
08-SEP-1999
15-SEP-1999
15-SEP-1999
05-OCT-1999
30-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                            cardiac hypertrophy; cardiovascular disorder; endothelial disorder; angiogenic disorder; atherosclerosis; osteoporosis; hypertension; myocardial infarction; diabetic retinopathy; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                            gene
                                                                                                                                                                                                                                                                                                                                                                                                                       Crohn's disease; psoriasis; endometriosis; ulcer; wound healing; c: Alzheimer's disease; Huntington's disease; stroke; drug screening;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human angiogenesis-associated protein PRO245, SEQ ID NO:91.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB53081;
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12-MAR-1999;
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            WPI; 2001-090793/10.
N-PSDB; AAC97437.
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                                                                                         (GETH ) GENENTECH INC.
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                                              NF,
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                                           J, Gurney AL,
Pitti RM, Wa
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                                                                                                                                                                                                                                                                                                                                                                                                            transgenic animal
                                                                     Baker KP,
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99WO-US21547
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                                              Watanabe CK,
                                                          Ferrara N,
Hillan KJ,
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                                               Williams PM,
                                                                     Gerber H,
                                                           Kuo SS,
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                                              Mark MR, M
                                                                     Goddard A;
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                                                           SA;
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Title:
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Gapop 10.0 , Gapext 0.5
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1637
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                                          219241 seqs, 76174552 residues
                                                                                                                                       1 MALRRPPRLRLCARLPDFFL.........VNYIRTDEEGDFRHKSSFVI 310
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        219241
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Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :

PIR_68:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	œ	7	S.	5)	S	2)		No.	Dog 11 t
154	154	154.5	155	155	155.5	155.5	159	159	159.5	159.5	160	160.5	160.5	161	161	161.5	. 163.5	164.5	169.5	169.5	•	171.5	174	177	177	180.5	186	187		Score	
9.4	9.4			9.5					9.7			9.8	9.8		9.8	9.9	10.0	10.0	10.4	•	10.5	10.5	10.6	10.8	10.8	•	11.4	11.4		Match 1	**
1863	1501	495	1907	344	1268	1259	4162	1323	6642	1033	7962	1612	761	1091	538	853	1051	3707	1912	858	1115	725	333	1092	725	1894	1088	725		Length	
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S46217	I58148	T25750	S50893	I56551	A39640	A43425	T42633	PN0568	T29757	S19247	I38346	T30805	IJHUNG	IJCHNL	JC2457	IJBONC	A39712	S18252	A56178	IJRTNC	IJMSNL	IJMSNG	A31923	JN0635	JE0100	C54689	IJXLNL	JE0099		ij	
protein-tyrosine-p	protein-tyrosine-p	hypothetical prote	protein-tyrosine-p		neural cell adhesi	Bravo/Nr-CAM cell	connectin/titin -	connectin 3B - chi	protein UNC-89 - C	cell adhesion prot		duttl protein - mo	cell	neural cell adhesi	vascular cell adhe		kinase-like protei	heparan sulfate pr	n-tyro	cell	cell	ë11	m prot	_	neural cell adhesi	protein-tyrosine-p	neural cell adhesi	neural cell adhesi		Description	

RESULT 2
IJXLNI.
IJXLNI.
IJXLNI.
IJXLNI.
IJXLNI.
N; Alternate names: NCAM-180
N; Alternate names: NCAM-180
N; Contains: neural cell adhesion molecule, short domain form (NCAM-140)
C; Species: Xenopus laevis (African clawed frog)
C; Decies: Nanopus laevis (African clawed frog)
C; Accession: S09600
R; Krieg, P.A.; Sakaguchi, D.S.; Kintner, C.R.

opioid-binding pro	JC1238	N	338	8.9	145.5	5
sax-3 protein - Ca	T42405	N	1273	8.9	146.5	4
biliary glycoprote	JC1509	N	458	8.9	146.5	ü
hypothetical prote	T29549	N	423	8.9	146.5	2
leukocyte antigen-	S46216	N	1898	9.0	147	Ë
leukocyte antigen-	TDHULK	ب	1897	9.0	147.5	ö
protein-tyrosine k	148697	<u>مبر</u>	881	9.0	147.5	39
protein-tyrosine k	148696	_	871	9.0	147.5	38
protein-tyrosine-p	A48758	_	1496	9.0	148	37
protein-tyrosine-p	B48758	_	1262	9.0	148	36
biliary glycoprote	S34338	N	521	9.1	149	35
transmembrane rece	T14160	N	1651	9.2	150	34
carcinoembryonic a	A36319	N	702	9.2	150.5	33
hypothetical prote	T17346	2	483	9.3	152	32
hypothetical prote	T33434	N	875	9.3	153	31
hypothetical prote	T33433	N	352	9.3	153	30

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ALIGNMENTS

Qy 198 NSETGTLVFTAVHKDDSGQYYCIASNDAGSA 228 :	Qy 138 KPVTPVCRVPKAVPVGKMATLHCQESEGH :: ::: Db 214 RQLRVNATAKMAESVVLSC-DADGF	Qy 83 KIQGDLAGRAEILGKTSLKIWNVTRRDSA : :	Qy 30 GAVNLKSNRTPVVQEFESVELSC	Query Match 11.4%; Score 187; DB 2; Length 729 Best Local Similarity 28.9%; Pred. No. 1.2e-07; Matches 61; Conservative 30; Mismatches 84; Indels	RESULT 1 JE0099 neural cell adhesion molecule 1 - African clawed frog N; Alternate names: N-CAM 1 C;Species: Xenopus laevis (African clawed frog) C;Date: 19-May-1998 #sequence_revision 29-May-1998 #text_cha A;Title: Molecular cloning of ssd-form neural cell adhesion A;Reference number: JE0099 A;Accession: JE0099 A;Acce
YCIASNDAGSA 228 : SCIANNQAGEA 291	KPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYSWYRNDVPLPTDSRANPRFRNSSFHL 197 : : :: :	83 KIQGDLAGRAEILGKTSLKIWNVTRRDSALYRCEVVARNDRKEIDEIV-IELTVQV 137 : : :	30 GAVNLKSSNRTPVVQEFESVELSCIITDSQTSDPR-IEWK-KIQDEQTTYVFFDN 82 	%; Score 187; DB 2; Length 725; %; Pred. No. 1.2e-07; 30; Mismatches 84; Indels 36; Gaps 11;	RESULT 1 JE0099 neural cell adhesion molecule 1 - African clawed frog N;Alternate names: N-CAM 1 C;Species: Xenopus laevis (African clawed frog) C;Date: 19 May-1998 #sequence_revision 29-May-1998 #text_change 21-Jul-2000 C;Accession: JE0099 C;Accession: JE0099 R;Kudo, M.; Takayama, E.; Tadakuma, T.; Shiokawa, K. Biochem. Biophys: Res. Commun. 245, 127-132, 1998 A;Title: Molecular cloning of ssd-form neural cell adhesion molecules (N-CAMs) as the A;Reference number: JE0099 A;Accession: JE0099 A;Accession: JE0099 A;Accession: JE0099 A;Accession: JE0099 A;Cross-references: DDBJ:AB008162; NID:93116226; PIDN:BAA25931.1; PID:93116227 A;Cross-references: DDBJ:AB008162; NID:93116226; PIDN:BAA25931.1; PID:93116227 A;Experimental source: heart C;Comment: This protein mediates and regulates various cell-cell interactions through C;Superfamily: neural cell adhesion molecule; fibronectin type III repeat homology; 1 F;413-475/Domain: immunoglobulin homology <imm> F;512-589/Domain: fibronectin type III repeat homology <imm <imm="" domain:="" f;512-589="" fibronectin="" h<="" homology="" iii="" repeat="" td="" type=""></imm></imm></imm></imm></imm></imm></imm></imm></imm></imm></imm></imm></imm></imm></imm></imm></imm>

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N;Contains: protein tyrosine phosphatase, receptor type delta, splice form C C;Species: Mus musculus (house mouse)
C;Date: 25-Apr-1995 #sequence_revision 19-May-1995 #text_change 12-Feb-1999
C;Accession: C54689; B54689
R;Mizuno, K.; Hasegawa, K.; Katagiri, T.; Ogimoto, M.; Ichikawa, T.; Yakura, Mol. Cell. Biol. 13, 5513-5523, 193
A;Title: MPTP delta, a putative murine homolog of HPTP delta, is expressed in A;Reference number: A54689; MUID:93360986
A;Accession: C54689
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F;618-679/Domain: fibronectin type III repeat homology <FN3B>
F;706-723/Domain: transmembrane #status predicted <TMM>
F;724-1088/Domain: intracellular #status predicted <TMM>
F;724-1088/Domain: intracellular #status predicted <TMM>
F;41-93,136-186,232-282,323-379,420-473/Disulfide bonds: #status predicted F;219,310,341,417,443,472/Binding site: carbohydrate (Asn) (covalent) #status
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A;Residues: 1-1088 <KRI>
A;Cesidues: 1-1088 <KRI>
A;Coss-references: EMBL:M25696; NID:g214609; PIDN:AAA49909.1;
A;Note: the authors translated the codon AAA for residue 970 as
C;Comment: NCAM mediates cell-cell adhesion via homophilic bind
C;Comment: Several forms of NCAM are produced by alternative sp
                                                                                                                                                                                                                                                                                                                                                                      protein-tyrosine-phosphatase (EC 3.1.3.48), receptor type delta, splice form B N,Alternate names: MFTP delta type B/C
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                       A; Experimental source: brain; splice form A; Nqte: sequence inconsistent with nucleot
                                                                        A; Molecule type: mRNA
A; Residues: 1-1894 <MIZ>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Gene: NCAM
                                                                                                                         A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic Acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; 20-8(
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Superfamily: neural cell adhesion molecule; fibronectin 
Keywords: alternative splicing; brain; cell adhesion; du
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Genetics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -705/Domain: extracellular #status predicted -95/Domain: immunoglobulin homology <IMMl>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         153/Region: 18-162/Region: 15-284/Domain: 381/Domain:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  475/Domain:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             188/Domain:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    198 NSETGTLVFTAVHKDDSGQYYCIASNDAGSA 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        )3,1050-1088/Product:
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sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYSWYRNDVPLPTDSRANPRFRNSSFHL 197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GTVNLKIYQKLTFKNAPTPQEFKEGEDAVIICDVSSSIPSIITWRHKGKD-----VIFKK 159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KIQGDLAGRAEILGKTSLKIWNVTRRDSALIRCE--VVARND--RKEIDEIV-IELTVQV 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DV-----RFVVLANNYLQIRGIKKTDEGTYRCEGRILARGEINYKDIQVIVNVPPTIQA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RQL----RVNATANMAESVVLSC-DADGFPDPEISWLKKGEPIEDG--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NEDQSEMTIHHVEKDDEAEYSCIANNQAGEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immunoglobulin homology <IMM2>
heparin binding #status predicted
heparin binding #status predicted
immunoglobulin homology <IMM3>
immunoglobulin homology <IMM4>
immunoglobulin homology <IMM4>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11.4%;
28.9%;
  from
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  NCBI
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Pred. No. 2.
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backbone
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.4e-07;
                         translation
(NCBIN:137486,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      molecule,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        84;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          long domain form #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 1088;
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duplication; heparin binding;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        binding with another ve splicing.
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NCBIP: 137487)
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                                                                                                                                                                                                 expressed in specialization
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-TPVVQEFESVELSCIITDSQTSDPR-IEWK-KIQDEQTTYVFFDN 82

Matches

Similarity

Conservative

28;

Mismatches

07:

88;

Indels

34;

Gaps

10;

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C;Accession: JE0100
C;Accession: JE0100
C;Accession: JE0100
R;Kudo, M.; Takayama, E.; Tadakuma, T.; Shiokawa, K
R;Kudo, M.; Takayama, E.; Tadakuma, T.; Shiokawa, K
Biochem. Biophys. Res. Commun. 245, 127-132, 1998
Biochem. Biophys. Res. Commun. 245, 127-132, 1998
A;Title: Molecular cloning of ssd-form neural cell
. nafarance number: JE0099; MUID:98204770
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            F;317-399/Domain: fibronectin type III repeat homology <FN3A> F;1278-1894/Domain: leukocyte common antigen cytosolic domain homology <I F;1278-1894/Domain: protein-tyrosine-phosphatase homology <PTP2> F;1536/Active site: Cys (phosphocysteine intermediate) #status predicted F;1542/Binding site: substrate phosphate (Arg) #status predicted F;1826/Active site: Cys (phosphocysteine intermediate) #status predicted F;1832/Binding site: substrate phosphate (Arg) #status predicted
                                                                                               C;Comment: This protein mediates and regulates various cell-cell interactions through C;Superfamily: neural cell adhesion molecule; fibronectin type III repeat homology; i F;413-475/Domain: immunoglobulin homology <IMM> F;512-589/Domain: fibronectin type III repeat homology <3FR>
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A;Note: sequence inconsistent with nucleotide translation
A;Note: sequence extracted from NCBI backbone (NCBIN:136527,
A;Note: sequence extracted from related protein; fibronecti
                                                                                                                                                                                                                                                             A; Molecule type: mRNA
A; Residues: 1-725 < KUD>
                                                                                                                                                                                                                                                                                                                 A; Reference number: JE0099;
A; Accession: JE0100
                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Xenopus laevis (African clawed frog)
C;Date: 19-May-1998 #sequence_revision 29-May-1998 #text_change
C;Accession: JE0100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        neural cell adhesion molecule 2 - N;Alternate names: N-CAM 2
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A; Residues: 1-352,'H',354-535,'S',537-601,1002-1894 <MI2>
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A; Status: preliminary
                                                                                                                                                                                                         A; Experimental source: heart
                                                                                                                                                                                                                                     A;Cross-references: DDBJ:AB008163; NID:g3116228; PIDN:BAA25932.1; PID:g3116229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JE0100
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10.8%;
Score 177;
Pred. No. 8
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Pred. No. 1.3e-06;
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DB.
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us-09-524-531a-15.rsp

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August 6, 2001, 09:38:58; Search time 13.3 Seconds (without alignments) 798.436 Million cell updates/sec
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1637
1 MALRRPPRLRLCARLPDFFL........VNYIRTDEEGDFRHKSSFVI 310
GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                 OM protein - protein search, using sw model
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BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

93435 seqs, 34255486 residues Searched:

93435 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% ... Maximum Match 100% Listing first 45 summaries

SwissProt_39:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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	Description	homo sapien	mus musculu	homo sapien	bos taurus	homo sapien	xenopus lae	xenopus lae	drosophila	mus musculu	mus musculu	rattus norv	homo sapien	rattus norv	homo sapien	mus musculu	mus musculu		gallus gall	mus musculu	bos taurus	homo sapien	homo			gallus gall		homo sapien	schistocerc	homo sapien	homo sapien	rattus norv	rattus norv	homo sapien
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ALIGNMENTS

RESULT	RESULT 1					
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ī	01-0CT-20	00 (Rel.	40, Last an	quence u notation	update)	
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8	Eukaryota; Metazoa;	; Metazoa			a; Vertebr	Craniata; Vertebrata; Euteleostomi;
ဗ	Mammalia;	Eutheria	; Primates;		ini; Homir	idae; Homo.
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1 2	TOWNER	220		DOTENTIAL	LULAR (FUI	ENITAD).
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FT	DISULFID	155	214	POTENTIAL.	Ľ.	
FT	CARBOHYD	98		N-LINKED) (POTENTIAL).
FT	CARBOHYD	187		N-LINKED) (POTENTIAL).
FT	CARBOHYD	236	236	ż	(GLCNAC.) (POTENTIAL).
SO	SEQUENCE	298 AA;			CA78E518E22DCAEE CRC64;	: CRC64;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     173 WYRNDVPLPTDSRANPRFRNSSFHLNSETGTLVFTAVHKDDSGQYYCIASNDAGSARCEE 232
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                                                                                                                                                         Gaps
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                    Length 298;
29.8%; Score 488; DB 1; Length 296
36.2%; Pred. No. 1.4e-34;
ive 60; Mismatches 115; Indels
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281 KATTMSENDFKHTKSFII 298
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superfamily that distributes at intercellular junctions and modulates monocyte transmigration."; MEDLINE-98327120; PubMed=9660867; Martin-Padura I., Lostaglio S., Schneemann M., Williams L., Romano Martin-Padura I., Lostaglio S., Schneemann M., Williams L., Romano Meruscella P., Panzeri C., Stoppacciaro A., Ruco L., Villa A., Simmons D., Dejana E.; C., Stoppacciaro A., Ruco L., Villa A., Jumons D., Dejana E.; C., Stoppacciaro A., Ruco L., Villa A., Junctional adheston molecule, a novel member of the immunoglobulin Junctional adheston molecule, a novel member of the immunoglobulin Mus musculus (Mouse). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 01-077-2000 (Rel. 40, Created) 01-0CT-2000 (Rel. 40, Last sequence update) 01-0CT-2000 (Rel. 40, Last annotation update JUNCTIONAL ADHESION MOLECULE PRECURSOR (JAM) 300 AA PRT; STANDARD; NCBI_TaxID=10090; JCAM OR JAM1. JAM1_MOUSE 088792; RESULT 2 JAM1_MOUSE

J. Cell Biol. 142:117-127(1998).
-1- FUNCTION: PLAYS A ROLE IN REGULATING MONOCYTE TRANSMIGRATION INVOLVED IN INTEGRITY OF EPITHELIAL BARRIER. INVOLVED IN PLATELET

SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN (POTENTIAL). TISSUE SPECIFICITY: LOCALIZED AT TIGHT JUNCTIONS OF BOTH EPITHELIAL AND ENDOTHELIAL CELLS.

SIMILARITY: CONTAINS 2 IMMUNOGLOBULIN-LIKE V-TYPE DOMAINS.

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EMBL; U89915; AAC32982.1; -. MGD; MGI:1321398; Jcam. IncerPro; IPR003006; -.

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 01-OCT-2000 (Rel. 40, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
JUNCTIONAL ADHESION MOLECULE PRECURSOR (JAM) (PLATELET ADHESION
MOLECULE 1) (PLATELET F11 RECEPTOR).
JCAM OR JAM1. 299 AA 01-OCT-2000 (Rel. 40, Created) STANDARD; Homo sapiens (Human). NCBI_TaxID=9606; JAM1_HUMAN Q9Y624; RESULT 3 JAM1_HUMAN

"Cutting edge: combined treatment of TNF-alpha and IFN-gamma causes redistribution of junctional adhesion molecule in human endothelial cells."; Ozaki H., Ishii K., Horiuchi H., Arai H., Kawamoto T., Okawa K., Iwamatsu A., Kita T.; SEQUENCE FROM N.A. MEDLINE-99323940; PubMed-10395639;

SEQUENCE FROM N.A. Sobocka M.B., Sobocki T., Rushbrook J.I., Banerjee P., Weiss C., Immunol. 163:553-557(1999). Kornecki E.;

"Molecular cloning and sequencing of the cDNA of Fl1 receptor, a novel Ig superfamily member from human platelets."; Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases.

Perfect score:

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Sequence:

Scoring table:

Searched:

Database

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Q9eqs8 mus musculu
Q10465 homo sapten
Q9ukv4 homo sapten
093242 gallus gall
Q2437 drosophila
099643 drosophila
                    Q9v3a5 drosophila
O02870 gallus gall
Q29123 sus scrofa
Q9tu80 canis famil
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091aj0 xenopus li
098918 gallus gal
057596 gallus gal
094915 gallus gal
09495 drosophila
094044 homo saplen
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mus musculu
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homo sapien
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        240
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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089026
Q9eqs9
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09tu80
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028939
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86.5%; Pred. No. 5.4e-129;
Live 18; Mismatches 24; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Curr. Top. Microbiol. Immunol. 251:91-98(2000).
EMBL; AJ300304; CAC20704.1; -.
SEQUENCE 310 AA; 34837 MW; 4B92BCB51D0A4B0A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Last sequence update)
Last annotation update)
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01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last seque
01-MAR-2001 (TrEMBLrel. 16, Last annot
JUNCTIONAL ADHESION MOLECULE-2, JAM-2
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O01761
Q08476
Q9IAJ0
Q98918
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Q10465
Q9UKV4
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089026
09EQS9
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Matches 268; Conservative
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NCBI_TaxID=10090;
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                                                                                                                                                                                                                                                                                                              1 MALRRPPRLRICARLPDFFL............310
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                                                                                                                                                                 Search time 35.5 Seconds
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                    GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                 425026 seqs, 132305027 residues
                                                                                                                                                                 6, 2001, 09:38:32;
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Maximum Match 100%
Listing first 45 summaries
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Q9JHY1
Q9JKD5
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Q9JKD5
Q9JKD5
Q9JKA2
Q9Z109
Q9JKA5
Q9PWR4
Q9
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Gapop 10.0 , Gapext 0.5
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1637
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sp_bacteria:*
sp_fungi:*
sp_human:*
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Maximum DB seq length: 200000000
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sp_rodent: *
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Match Length
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Query Match
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6
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01-0CT-2000 (TrEMBLrel. 15, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
VASCULAR ENDOTHELIAL JUNCTION-ASSOCIATED MOLECULE (JUNCTIONAL ADHESION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Aurrand-Lions M.A., Duncan L., Du Pasquier L., Imhof B.A.; "Cloning of JAM-2 and JAM-3: an Emerging Junctional Adhesion Molecular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    φ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Palmeri D., van Zante A., Huang C.-C., Hemmerich S., Rosen S.D.; vsscular Endothelial Junction-associated Molecule, a Novel Member the Immunoglobulin Superfamily, is Localized to Intercellular Boundaries of Endothelial Cells.";
                                                                 2 ARSPQGLIALLILLHYLIVALDYHKANGFSASKDHRQEVTVIEFQEAILAC-KTPKKTTSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutharia; Rodentia; Sciurognathi; Muridae; Murinae; Mus
NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13 ARLPDFFLLLLFRGCLIGAVNLKSSN------RTPVVQEFESVELSCIITDSQTSDP
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                                            241 NIGGIIGGVLVVLAVLALITLGICCAYRRGYFINNKQDGESYKNPGKPDGVNYIRTDEEG
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36.7%; Pred. No. 9.3e-39;
ive 54; Mismatches 116; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      298 AA; 33047 MW; 1124E0F07E6CF751 CRC64;
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EMBL; AF255911; AF81224.1; -.
EMBL; AF2157; CAC20699.1; -.
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Matches 114; Conservative
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SMART; SM00408; IGC
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Rattus norvegicus (Rat).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
NCBI_TaxID-10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TTYVFFDNKIQGDLAGRAEILGKTSLKIWNVTRRDSALYRCEVVARNDRKEIDEIVIELT 134
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28.3%; Score 463.5; DB 11; Length 300; 37.9%; Pred. No. 1.6e-36; Live 49; Mismatches 121; Indels 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mashima H., Koʻlima I.;
Submitted (JUN-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AR276998; AR78250.1; -.
InterPro; IPR003006; -.
InterPro; IPR003066; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AMART: PRO0047; 1g; 2.
SMART: SMO0406; IGv; 1.
SFROUENCE 300 AA; 32369 MW; 45AE362A96158BFA CRC64;
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Catarrhini; Hominidae;
                                                                                                                                                                                                                       01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
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Mammalia; Eutheria; Primates;
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Foley C., Parkos C.A.,
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Copyright (c) 1993 - 2000 Compugen Ltd
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Seguence 19, Appl	Seguence 21, Appl	Sequence 56, Appl	Sequence 48, Appl	Sequence 44, Appl	Seguence 46, Appl	Sequence 2, Appl1	Sequence 41, Appl	Sequence 31, Appl	Sequence 1, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 26, Appl	Sequence 2, Appl1	25,	Sequence 2, Appli	Sequence 5, Appl1	Sequence 12, Appl	•
PCT-US95-08493-19	PCT-US95-08493-21	US-08-414-657D-56	US-08-414-657D-48	US-08-414-657D-44	US-08-414-657D-46	US-08-414-657D-2	US-08-414-657D-41	US-08-408-095-31	US-08-374-834-1	US-08-644-271-1	US-08-217-299-1	US-08-928-383B-26	US-09-040-774-2	US-09-041-886-25	PCT-US94-05277-2	US-08-986-485-5	US-08-752-307B-12	
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145.5	145.5	144.5	144.5	144.5	144.5	144.5	144.5	144	143.5	143.5	143	142	140.5	138.5	138.5	138	137	
28	58	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	

ALIGNMENTS

127 120 187 180 246 240 240 302

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APPLICANT: Old, Lloyd J.; Welt, Sydney; Ritter, Gerd;
APPLICANT: Simpson, Richard J.; Nice, Edouard; Moritz, R. L.;
APPLICANT: Catimel, B.; Ji, Hong; Burgess, Anthony W.;
APPLICANT: Heath, Joan K.; White, Sara J.; Johnstone, Cameron JITLE OF INVENTION: Colon Cell And Colon Cancer Cell
TITLE OF INVENTION: Associated Nucleic Acid Molecules, Protein And Peptides NUMBER OF SEQUENCES: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          63 KFDQGDTTRLVCYNNKITASYEDRVTFL-PTGITFKSVTREDTGTYTC-MVSEEGGNSYG 120
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                                                                                                                                                                                                                                                     APPLICANT: Ortust, Recent APPLICANT: Outset, Recent APPLICANT: Outset, Recent APPLICANT: Murison, James Greg TITLE OF INVENTION: Compositions Isolated From Skin Cells TITLE OF INVENTION: and Methods For Their Use FILE REFERENCE: 11000.1011c1 CURRENT APPLICATION NUMBER: US/09/188,930A CURRENT FILING DATE: 1998-11-09 NUMBER OF SEQ ID NOS: 348 SEQ ID NOS: 348 SEQ ID NOS: 348 LENGTH: 299
                                                                                                                ; Sequence 189, Application US/09188930A
; Patent No. 6150502
; GENERAL INFORMATION:
; APPLICANT: Watson, James D.
; APPLICANT: Strachan, Lorna
; APPLICANT: Sleeman, Matthew
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Sequence 22, Application US/08597495B

Patent No. 2712369

GENERAL INFORMATION:

APPLICANT: Old, Lloyd J.; Welt, 8
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Matches 99; Conservative
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; NAME/KEY: UNSURE
; LOCATION: (289)...(289)
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292 KQTSSFLV 299
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292 KQTSSFLV 299
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ORGANISM: Human
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                                                                                                                                                                                                       Diskette, 3.5 inch, 360 kb storage
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14.0%; Score 228.5; DB 1;
Best Local Similarity 25.7%; Pred. No. 2.4e-15;
Matches 79; Conservative 49; Mismatches 128;
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Patent No. 5942606

GENERAL INFORMATION:
APPLICANT: Lal, Preeti
APPLICANT: Corley, Neil C.
TITLE OF INVENTION: VIRAL RECEPTOR PROTEIN
UMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/597,495B FILING DATE: 02-Feb-1996 CLASSIFICATION: 435 PRIOR APPLICATION: 435 PRIOR APPLICATION DATE: APPLICATION NUMBER: 08/511,876 PTLING DATE: 04-Aug-1995 ATTORNEY/AGENT INFORMATION: NAME: Hanson, No. 5712369man D. REGISTRATION NUMBER: 30,946 REFERENCE/DOCKET NUMBER: LUD 5316.1 TELECOMMUNICATION INFORMATION:
Felfe & Lynch
LAGET: 805 Third Avenue
CITY: New York
STATE: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (212) 688-9200
TELEFAX: (212) 838-3884
INFORMATION FOR SEQ ID NO: 22
SEQUENCE CHARACTERISTICS:
LENGTH: 319 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                       PC-DOS
                                                                                                                                                                                                                                                                              SOFTWARE: Wordperfect CURRENT APPLICATION DATA:
                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                           COMPUTER: IBM PS/2
OPERATING SYSTEM: I
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Listing first 45 summaries
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Copyright (c) 1993 - 2000 Compugen
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Human confluency
Human PRO1868 pro
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Human PRO301 prote	AAB80232	22	299	25.4	415	ر.
Amino acid sequenc	AAB31202	22	299	25.4	415	4
Human A33 receptor	AAY76076	21	299	25.4	415	w
rec	AAY76011	21	299	25.4	415	N
Human PRO301 prote	AAY70670	21	299	•	415	_
Human PRO301 antit	AAY95344	. 21	299	•	415	0
	AAB24405	21	299	•	415	9
_	AAW74464	20	299	25.4	415	8
Amino acid sequenc	AAY13364	20	299	25.4	415	7
Amino acid sequenc	AAY23321	20	299	25.4	415	Ó
PRO307 prot	AAY08071	20	299		415	S
Human confluency r	AAB27277	21	213	•	425	4
gen	AAB70500	22	215	٠	428	ω
Gene 15 human secr	AAB39253	21	280	27.3	445	Ň
A33 related antige	AAY23325	20	300	•	449.5	Ĺ
Mouse junctional a	AAW61380	19	300		449.5	Ò
Human PRO245 prote	AAB50904	22	312		474	Ö
w	AAB53081	22	312		478	œ
PRO245	AAB80222	22	312	29.3	478	7
	AAY70668	21	312		478	6
PRO245	AAB24401	21	312		478	ű
Human PRO245 prote	AAB33421	21	312		478	4
Amino acid sequenc	AAY13354	20	312	29.3	478	ũ
elated a	AAY23324	20	312	٠	478	ຄ
PRO245	AAY08060	20	312	•	478	ï
æ	AAB27275	21	298		490.5	Õ
Human confluency r	AAB27273	21	298		490.5	٥
Human secreted pro	AAW75220	19	298		498	8
ů	AAU00512	22	298	30.9	505	7
ted protei	AAW85457	19	298		505	Ò
secreted	AAB39254	21	285	٠	1313	'n
1G	AAY96294	21	310	85.8	1401	4
≱13 associ	AAB80431	22	339	86.3		w
Human secreted pro	AAB38384	21	311	86.3	1409	N

ALIGNMENTS

RESULT AAB27272

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AAB27272 standard; Protein;

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23-FEB-2001 AAB27272;

(first

entry)

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Isolated human Confluency Regulated Adhesion Molecule 1 or 2 (CRAM-1 or CRAM-2) polypeptide, useful for treatment of tumors, inflammation reactions and modulating vascular permeability -
                                                                                                                                                                                                                                 Immunoglobulin superfamily; Ig Sf; vascular adhesion molecule; inflammation; cancer; wound; angiogenesis; human; confluency regulated adhesion molecule 1; CRAM-1; JAM-2.
                                                                          Imhof BA,
                                                                                               (RMFD-) RMF DICTAGENE
                                                                                                                      11-MAR-1999;
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                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                               Human confluency regulated adhesion molecule 1 #1.
                                                     2000-587436/55
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Isolated human Confluency Regulated Adhesion Molecule 1 or
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                                                                  (RMFD-) RMF DICTAGENE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      {\tt CRAM-2)} polypeptide, useful for treatment of tumors, inflammation reactions and modulating vascular permeability -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                              WO200053749-A2
                                                                                                                                                                                                       Immunoglobulin superfamily; Ig Sf; vascular adhesion molecule; inflammation; cancer; wound; angiogenesis; human;
                                                                                                                                                                                                                                           Human confluency regulated
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                                                                                                                                                                                           regulated
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 1633; DB 21; ilarity 100.0%; Pred. No. 3.6e-124; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                   (first entry)
                                                                       99EP-0200746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                59pp; English
                                                                                                                                                                                                                                                                                                                     Protein;
                                                                                                                                                                                             adhesion molecule
                                                                                                                                                                                                                                             adhesion
                                                                                                                                                                                                                                             molecule 1 #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ļength
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WPI; 2000-587436/55

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                                                           osteoarthritis; spondyloarthropathy; systemic sclerosis; sarcoidosis; idiopathic inflammatory myopathy; Sjogren's syndrome; thyroiditis; systemic vasculitis; autoimmune haemolytic anaemia; diabetes mellitus; autoimmune themolytic anaemia; diabetes mellitus; autoimmune thrombocytopaenia; immune-mediated renal disease; autoimmune thrombocytopaenia; immune-mediated; inflammatory bowel disease; hepatobiliary disease; whipple's disease; inflammatory bowel disease; gluten-sensitive enteropathy; autoimmune disease; immune-mediated skin disease; allergic disease; autoimmune disease; immune-mediated skin disease; allergic disease;
                                                                                                                                                                        Human; immune related disease; diagnosis; antiinflammatory; cardian dermatological; antiarthritic; antiheumatic; immunosuppressive; haemostatic; antithyroid; antidiabetic; nootropic; neuroprotective; antianaemic; hepatotropic; virucide; antipsoriatic; antiallergic; antiasthmatic; systemic lupus erythematosus; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is the human confluency regulated adhesion molecule 1 (CRAM-1, also known as JAM-2). CRAM-1 is one of the vascular adhesion proteins of the immunoglobulin superfamily (Ig Sf). The CRAM-1 protein and coding sequence can be used in the treatment of cancer, inflammation, to modulate cell-cell interactions and angiogenesis, and in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated human Confluency Regulated Adhesion Molecule 1 or 2 (CRAM-1) or CRAM-2) polypeptide, useful for treatment of tumors, inflammation reactions and modulating vascular permeability -
                                                                                                                                                                                                                                                                              Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ches
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                              immunological disease;
graft rejection; graft.
                                                                                                                                                                                                                                                                                                                                                                                              157
                                                                                                                                                                                                                                                                                                                                                                                                                                                             301
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                                                                                                                                                                                                                                                                                                                                                                             AB33457 standard; Protein; 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61
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Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        d coding sequence can be u
modulate cell-cell intera
dulation of wound healing.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             dfrhkssfvi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        niggiiggvlvvlavlalitlgiccayrrgyfinnkqdgesyknpgkpdgvnyirtdeeg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NIAGIIGGVLVVLIVLAVITMGICCAYRRGCFISSKQDGESYKSPGKHDGVNYIRTSEEG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ndrkeideivieltvqvkpvtpvcrvpkavpvgkmatlhcqeseghprphyswyrndvpl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NDRKEYDEITIELIVQVKPYTPYCRIPAAVPYGKTATLQCQESEGYPRPHYSWYRNDYPL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SDPRIEWKKIQDGQTTYVYFDNKIQGDLAGRTDVFGKTSLRIWNVTRSDSAIYRCEVVAL 120
                                                                                                                                                                                                                                                                              PRO1868
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           310 AA;
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                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6; 59pp; English
                                                                                                                                                                                                                                                                          protein UNQ859
                              graft-versus-host-disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            85.8%;
                                               transplantation associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 1409; DB 21;
Pred. No. 4.5e-106;
8; Mismatches 26;
                                                                                                                                                                                                                                                                            SEQ ID
                                                                                                                                                                                                                                                                            NO:193.
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12-AFR-1999
20-AFR-1999
21-AFR-1999
21-AFR-1999
21-JUN-1999
22-JUN-1999
23-JUN-1999
23-JUN-1999
23-JUL-1999
26-JUL-1999
26-JUL-1999
27-JUL-1999
28-JUL-1999
28-JUL-1999
29-CCT-1999
15-SEP-1999
15-SEP-1999
15-SEP-1999
15-SEP-1999
15-SEP-1999
15-DCT-1999
29-NOV-1999
29-NOV-1999
29-NOV-1999
29-DEC-1999
20-DEC-1999
20-DEC-1999
21-DEC-1999
                     The present invention describes sixty four human PRO proteins which can be used in the treatment of immune related diseases. The human PRO proteins, anti-PRO antibodies, agonists and antagonists are useful for treating and diagnosing immune related disorders. The disorders are selected from systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, juvenile chronic arthritis, spondyloarthropathies, systemic sclerosis, idiopathic inflammatory myopathies, Sjogren's systemic vasculitis, sarcoidosis, autoimmune haemolytic anamemia, autoimmune thrombocytopaenia, thyroiditis, diabetes mellitus, anamemia, autoimmune thrombocytopaenia, thyroiditis, diabetes mellitus,
                                                                                                                                                                                           immune rel
arthritis,
                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                             Stewart
                                                                                                                                                                                                                                                                                                          Kabakoff
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-MAR-1999;
12-MAR-1999;
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             immune-mediated renal disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-MAR-1999;
                                                                                                                                                                                                                                                   2000-572271/53.
DB; AAC58622.
                                                                                                                                                                                                                       four PRO
                                                                                                                                                                                                          related
                                                                                                                                                                                                                                                                                                                                                   GENENTECH INC.
                                                                                                                                                                                                                                                                                                        RC,
                                                                                                                                                                Fig 88;
                                                                                                                                                                                             osteoarthritis,
                                                                                                                                                                                                                                                                                                                                                                           99US-0128849.
99WO-US08615.
99US-0131445.
99US-0131252.
99US-0141037.
99US-01445698.
99US-01445698.
99US-0146222.
99WO-US20594.
99WO-US20594.
99WO-US21547.
99WO-US21547.
99WO-US21547.
99WO-US21630.
99WO-US21630.
99WO-US21831.
99WO-US28301.
99WO-US28301.
99WO-US28301.
99WO-US2855.
                                                                                                                                                                                                                                                                                           Tumas D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000WO-US05841
                                                                                                                                                                                                                                                                                                        Lu Y,
                                                                                                                                                                                                     polypeptides, useful in the diagnosis and treatment of disorders, e.g. systemic lupus erythematosis, rheumato
nervous systems,
                                                                                                                                                                                                                                                                                                                       Baker
                                                                                                                                                                                                                                                                                                          Pan
                                                                                                                                                                                                                                                                                                                        ΚP,
                                                                                                                                                                                                                                                                                           an J, Pennica
Watanabe CK,
                                                                                                                                                                English.
                                                                                                                                                                                                                                                                                                                       Goddard A,
                                                                                                                                                                                           e.g. systemic lupus erythematosis, thyroiditis and diabetes mellitus
             demyelinating diseases of the central
hepatobiliary diseases,
                                                                                                                                                                                                                                                                                Gurney AL, ... S, Shelton DL, S'
                                                                                                                                                                                                                                                                                           Wood WI,
                                                                                                                                                                                                                                                                                                          Hebert C,
DL, Smith
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                  Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           bowel disease, gluten-sensitive enteropathy and Whipple's disease, autoimmune or immune-mediated skin diseases, allergic diseases, immunological diseases of the lung, and transplantation associated diseases including graft rejection and graft-versus-host-disease. AAC5837 to AAC58578 represent PCR primers and hybridisation probes in the isolation of human PRO sequences. AAC58579 to AAC58642 and AAB33414 to AAB33477 represent human PRO polynucleotide and protein sequences given in the exemplification of the present invention.
Modified-site
                                                                                                                                               Modified-site
                                                                                                                                                                                Key
                                                                                                                                                                                                                                                                                                       AAY96735;
                                                                                                                                                                                                                                                                                                                         AAY96735 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                         Modified-site
                                                                             Modified-site
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                                                                                                                          Modified-site
                                                                                                                                                                    Peptide
                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                          PRO1868, an A33
                                                                                                                                                                                                                                                                                  6-SEP-2000
                                                                                                                                                                                                                           nti-inflammatory;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 MALSRRLRLRLYARLPHFFLLLLFRGCMIEAVNLKSSNRNPVVHEFESVELSCIITHSQT 60
                                                                                                                                                                                                                                                                                                                                                  υ
                                                                                                                                                                                                                                                                                                                                                                                   dfrhkssfvi 310
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NDRKEYDEITIELIVQVKPYTPYCRIPAAYPYGKTATLQCQESEGYPRPHYSWYRNDYPL 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DFRHKSSFVI 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PTDSRANPRFQNSSFHVNSETGTLVFNAVHKDDSGQYYCIASNDAGAARCEGQDMEVYDL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    266;
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                                                                                                                                                                                                                                      antigen;
                                                                                                                                                                                                                                                                                (first entry)
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                   phosphorylation 192..195
  /note=
215..2
                                                                                                                                                                                                                                                           antigen homologue.
                                                        /note- "(
                                                                                                                         /note= "N-myristoylation site"
69..77
                                                                                                                                               /label= Signal_peptide
26..31
                                                                              106..109
                                                                                                    104..107
                                                                                                                                                                              Location/Qualifiers
                                                                                        note-
                                                                                                            note- "Tyrosine kinase phosphorylation
                                                                                                                                                                                                                        en; secreted protein; transmembrane protein
cytostatic; recombinant production; gene t)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             85.8%;
                                            "CAMP- and cGMP-dependent protein kinase
                                                                                        "N-glycosylation site"
            "N-glycosylation site
                                                                  "Casein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18;
                                                                                                                                                                                                                                                                                                                            310 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 1409; DB 21;
Pred. No. 4.5e-106;
L8; Mismatches 26;
                                                                    kinase
                                                                 II phosphorylation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                          gene therapy
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                                                                    site"
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                                                                                                                       Matches 266;
                                                                                                                                    Query Match
Best Local
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09-FEB-1999;
10-FEB-1999;
12-FEB-1999;
02-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Botstein D,
Gurney AL,
                                                                                                                                                                                                                             This is PRO1868, a putative homologue of A33 antigen, a known colorectal cancer-associated marker. The invention concerns novel secreted and transmembrane proteins, designated PRO polypeptides. The CDNA and gene sequences are useful in the recombinant production of PRO polypeptides, as a hybridization probe to screen libraries to isolate CDNAs with sequence identity to PRO polypeptides or to map the gene encoding the PRO polypeptides and analyzing genetic disorders. The CDNA/gene can also be used to produce transgenic animals useful for the
                                                                                                                                                                                 Sequence
                                                                                                                                                                                                       development and screening of the
rapeutically useful reagents. also be used in gene the
rapy, e.g. to replace a defective gene
                                                                                                                                                                                                                                                                                                                                          Claim 1; Fig 14; 154pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                      N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-DEC-1998;
22-DEC-1998;
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12-JAN-1999;
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 121
                                                                                                                                                                                                                                                                                                                                                                             solated nucleic acid molecule encodes a PRO polypeptide which
                         61
                                               61
                                                                                                                                    Local Similarity
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                                   SDPRIEWKKIQDGQTTYVYFDNKIQGDLAGRTDVFGKTSLRIWNVTRSDSAIYRCEVVAL 120
                                                                      2000-431586/37.
DB; AAA51265.
  NDRKEVDEITIELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWYRNDVPL
                       sdpriewkkiqdeqttyvffdnkiqgdlagraeilgktslkiwnvtrrdsalyrcevvar
                                                                                              MALSRRLRLYARLPHFFLLLLFRGCMIEAVNLKSSNRNPVVHEFESVELSCIITHSQT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                ָס,
                                                                                                                                                                                 310 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pan
                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Desnoyers L,
an J, Roy MA,
                                                                                                                                                                                                                                                                                                                                                                polypeptide
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99US-0119537.
99US-0119965.
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99US-0115565.
99US-0115733.
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296..299
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243..263
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226..231
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, Stewart TA,
                                                                                                                                  Score 1409; DB 21; Pred. No. 4.5e-106;
                                                                                                                       Mismatches
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A, Tumas
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Watanabe CK;
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                                                                                                            Ashkenazi AJ, Botst
Filvaroff E, Fong S
Godowski PJ, Grimal
Mather JP, Pan J,
                                                                                                                                                                                    13-SEP-1999
15-SEP-1999
15-SEP-1999
05-OCT-1999
05-OCT-1999
29-NOV-1999
16-DEC-1999
20-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                       Human; PRO; dermatological; antipsoriatic; cytostatic; antiinflammatory; antiparkinsonian nootropic; neuroprotective; vulnerary; cardiant; antiangiogenic; vasotropic; antiasthmatic; antiheumatic; cancer; antiarthritic; antiinfertility; antidiabetic; antiviral; diabetes; ophthalmological; gene therapy; skin disease; gastrointestinal disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7272
                           Alzheimer's
                                   the treatment of skin disease squamous cell carcinoma) and
                                                                                                                                                                                                                                                                         08-SEP-1999;
                                                                                                                                                                                                                                                                                  26-JUL-1999;
28-JUL-1999;
                                                                                                                                                                                                                                                                                                   07-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB80272 standard; Protein; 310 AA.
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DB; AAF72433.
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                                             one nucleic acids encoding PRO reatment of skin diseases (e.g.
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                           disease)
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                                                                                                            Botstein D, Desnoyers L,
Fong S, Gao W, Gerber H,
Grimaldi CJ, Gurney AL, H
an J, Paoni NF, Roy MA, S
                                                                                                    Wood WI;
                                                                                                                                                                            99WO-US20944.
99WO-US21090.
99WO-US21347.
99WO-US23089.
99WO-US28214.
99WO-US30095.
99WO-US30915.
99WO-US30999.
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99US-0145698.
99US-0146222.
99WO-US20594.
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                                  ding PRO polypeptides which are useful in es (e.g. psoriasis), cancers (e.g. lung neurodegenerative diseases (e.g.
                                                                                                            Hillan KJ,
Stewart TA,
                                                                                                             Kljavin
Tumas [
                                                                                                                               Goddard
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Best Local :
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         Rosen
Shi Y,
                                                                                                                                                                 Secreted protein; human; autoimmune; hyperproliferation; cardiovascular; cerebrovascular; infection; food.
                                                                                                      01-FEB-2001.
                                                                                                                          WO200107459-A1
                                                                                                                                                                                                 Secreted
                                        (HUMA-) HUMAN GENOME
                                                                                 20-JUL-2000;
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                                                                                                                                                                                                                                                              AAB80383 standard; protein;
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                                                                                                                                                                                                                      (first entry)
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Olsen HS,
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Pred. No. 4
          Duan RD,
Birse CE,
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4.5e-106;
hes 26;
                   Ni J,
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                      23-JUL-1999;
                                                                    20-JUL-2000; 2000WO-US19735
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                                                                                                                                                                                                                                                                     Secreted protein; cardiovascular; c
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ABB0408 standard; protein; 310 AA.
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                      99US-0145220
                                                                                                                                                                                                                                                                     cerebrovascular; infection;
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                                                                                                                                                                                                                                                                                         human; autoimmune; hyperproliferation;
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Pred. No. 4.5e-106;
18; Mismatches 26;
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                                                                                                                                                   Secreted protein; cardiovascular; co
                                                                                                                                                                                                                                                                                                                                                                                   AAB80409 standard; protein;
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                                                                                                    Homo sapiens.
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, Lafleur DW,
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                                                                                                                                                      cerebrovascular;
                                                                                                                                                      human; autoimmune; hyperproliferation;
erebrovascular; infection; food.
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Olsen HS,
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Pred. No. 4.5e-106;
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Komatsoulis GA;
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Best Local S
Matches 266
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Immunosuppressive; antiarthritic; au
cytostatic; cardiant; vasotropic; co
nootropic; antibacterial; virucide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New isolated nucleic acid encoding 29 secreted proteins, for diagnosing, preventing and treating e.g. autoimmune, hyperproliferative, cardiovascular, and ocular diseases or disorders
                                           Human secreted
                                                                 31-JAN-2001
                                                                                                        AAB38333 standard; Protein;
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                                                                                                                                                                                    DFRHKSSFVI 310
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                                         protein encoded by gene 13 clone HAPSA79.
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Pred. No. 4.5e-106;
8; Mismatches 26;
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Birse CE,
antirheumatic; antiproliferative;
cerebroprotective; neuroprotective;
e; fungicide; opthalmalogical; human;
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                                                                                                                                                                                                                                                                                            Matches
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Best Local Similarity
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Lafleur DW,
                                                                                                                                                                                                                                                                                                                                                                  Sequence
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26-APR-1999;
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                                                                                                                 ndrkeideivieltvqvkpvtpvcrvpkavpvgkmatlhcqeseghprphyswyrndvpl
                                                         ptdsranprfrnssfhlnsetgtlvftavhkddsggyyciasndagsarceegemevydl
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99US-0130991.
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Pred. No. 4.5e-106;
.8; Mismatches 26;
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                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                         disorders e.g.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New nucleic acid molecules encoding 62 human secreted proteins for diagnosing, preventing, treating or ameliorating medical conditions used as food additives or preservatives -
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26-APR-1999;
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                                                                                                                                                                                                                                                                              orders e.g. corneal infection. The polypeptides can also be used to wound healing and epithelial cell proliferation, to prevent skin ng due to sunburn, to maintain organs before transplantation, for porting cell culture of primary tissues, to regenerate tissues and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000-647418/62
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Olsen HS, Ebner
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                                                                                                                86.3%;
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                                                                                            Score 1409; DB 21;
Pred. No. 4.5e-106;
8; Mismatches 26;
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         Sequences AAB38321-B38396 represent the amino acid sequences of 62 human secreted proteins encoded by the genes AAC69512-C6587. The genes and proteins are useful for preventing, ameliorating or treating medical conditions, e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant)agonists are useful in the diagnosis, treatment and prevention of: (a) autoimmune diseases e.g. rheumatoid arthritis; (b) hyperproliferative disorders e.g. neoplasms of the breast or liver; (c) cardiovascular disorders e.g. cardiac arrest; (d) cerebrovascular disorders e.g. crebral ischemia; (e) angiogenesis; (f) nervous system disorders e.g. Alzheimer's disease; (g)
                                                                                                                                                                                                                               New nucleic acid molecules encoding 62 human secreted proteins for diagnosing, preventing, treating or ameliorating medical conditions and used as food additives or preservatives -
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Lafleur DW,
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26-APR-1999;
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               New isolated nucleic acid encoding 29 secreted proteins, for diagnosing, preventing and treating e.g. autoimmune, hyperproliferative, cardiovascular, and ocular diseases or disorders
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scular; cerebrovascular; infection; food.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           dfrhkssfvi 339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DFRHKSSFVI 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NDRKEVDEITIELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWYRNDVPL 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ndrkeideivieltvqvkpvtpvcrvpkavpvgkmatlhcqeseghprphyswyrndvpl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IGFAM-6
                                                                                                                                                                                                                                                                                                                                                                                                                            immunoglobulin; IGFAM-6; IGFAM; immune disorder;
                                                                                                                                                                                                                                                                                                                                                                                                    inflammation; haematopoiesis; AIDS; allergy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 75;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunoglobulin.
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                                                                                                                                                                                                                                             /label=
31..310
                                                                                                         /label=
238..260
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/label=
                                                                                                                                                         153..221
                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                   /label=
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85.8%;
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Pred. No. 5e-106;
18; Mismatches 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              English
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                                                                                                                                                                                                                                                                                                                                                                                                                                cancer;
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                                          RESULT
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Best Local Sim
Matches 265;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             multiple sclerosis, psoriasis, rheumatoru architich, complications systemic lupus erythematosus and ulcerative colitis), complications cancer, haemodialysis and extracorporeal circulation, trauma and haematopoietic cancer (such as leukaemia) and infections caused by bacteria, viruses, fungi or parasites.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 arteriosclerosis, asthma, atherosclerosis, cholecystitis, Crohn's disease, diabetes mellitus, emphysema, Graves' disease, hepatitis, multiple sclerosis, psoriasis, rheumatoid arthritis, scleroderma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yue
              AAB39254 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cardiovascular tissue, where cancer and inflammation are common. The gene, protein, its antibodies, agonists and antagonists are suitable for diagnosing and treating many diseases, including cancer, immune system disorders (such as inflammation, AIDS, allergies, anaemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunoglobulin superfamily proteins, the agonist and antagonist of the protein is useful for preventing and treating disorders associated with altered levels of the protein such as cancer, immune system disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (INCY-) INCYTE PHARM INC
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22-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GFAM-6. Its gene was isolated from a cDNA library of clasue. It is expressed in reproductive, nervous and
                                                                                    301
                                                                                                                                                                        241
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                                                                                                                                                                                                                                                                                                                                                                    DAM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present sequence
                                                                                     dfrhkssfvi 310
                                                                                                                                          NIAGIIGGVLVVLIVLAVITMGICCAYRRGCFISSKQDGESYKSPGKHDGVNYIRTSEEG
                                                                                                                                                                                                                                                                      NDRKEVDEITIELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWYRNDVPL 180
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)B; AAA27386.
                                                                                                                DFRHKSSFVI 310
                                                                                                                                                                                                  ptdsranprfrnssshlnsetgtlvftavhkddsggyyciasndagsarceeqemevydl
                                                                                                                                                                                                                                                       ndrkeideivieltvqvkpvtpvcrvpkavpvgkmatlhcqeseghprphyswyrndvpl
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                                                                                                                                                                                                                                                                                                                                           SDPRIEWKKIQDGQTTYVYFDNKIQGDLAGRTDVFGKTSLRIWNVTRSDSAIYRCEVVAL 120
                                                                                                                                                                                                                             PTDSRANPRFQNSSFHVNSETGTLVFNAVHKDDSGQYYCIASNDAGAARCEGQDMEVYDL
                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
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98US-0113635.
99US-0128194.
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Hillman
                                                                                                                                                                                                                                                                                                                                                                                                                                              85.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  is the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 is the human immunoglobulin superfamily protein isolated from a cDNA library of leg
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                                                                                                                                                                                                                                                                                                                                                                                                                               18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 1401;
Pred. No. 2
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Yang J;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                              2e-105;
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86

Matches

Similarity

Conservative

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Mismatches No.

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26 ,...

GCMIEAVNLKSSNRNPVVHEFESVELSCIITHSQTSDPRIEWKKIQDGQTTYVYFDNKIQ 85

GDLAGRTDVFGKTSLRIWNVTRSDSAIYRCEVVALNDRKEVDEITIELIVQVKPVTPVCR 145 gcligavnlkssnrtpvvgefesvelsciitdsqtsdpriewkkiqdeqttyvffdnkiq 60

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DRX PRACE TO A CONTRACT OF A C
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                                                                                                                                                                                                                                                                                        cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; and opthalmalogical. The human secreted proteins, polynucleotides, antagonists and agonists of the invention may be useful in the treatment, prevention, and/or diagnosis of various disease, disorders and conditions such as autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cardiac arrest, verebrovascular disorders e.g. cardiac arrest, verebrovascular disorders e.g. cerebroliferation in the polypeptides and disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to regenerate tissues, maintain organs before transplantation, in chemotaxis and as a food additive or preservative e.g. to increase storage capabilities. Sequences AAC74214-C74222 and AB39178 are used and also the increase distribution and absorber the increase of the increase and an antibal properties of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           secreted proteins represented in AAB39179-B39226. Sequences AAB39227-B39308 are alternative proteins encoded by the genes, and als protein sequences with which they share homology. The proteins have activities based on the tissues and cells in which they are expressed. Examples of activities include: immunosuppressive; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       opthalmalogical; autoimmune disease; rheumatoid arthritis; angiogenesis; hyperproliferative disorder; cardiovascular disorder; infection; cerebrovascular disorder; nervous system disorder; ocular disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; secreted protein; immunosuppressive; antiarthy antiproliferative; cytostatic; cardiant; vasotropic; nootropic; neuroprotective; antibacterial; virucide;
                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 32; 434pp; English.
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10-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isolated nucleic acid molecule encoding a human secreted protein used in preventing, treating or ameliorating a medical condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2000-579483/54.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ruben SM,
                                                                                                                                                                                                                                                                 isolation
                                                                                                                                                                           285 AA;
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99US-0169980.
                                                                                                                                                                                                                                                                 and
80.4%;
85.6%;
                                                                                                                                                                                                                                                                 characterisation
                                        Score 1313;
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suppressing activity; haematopolesis regulating activity; tissue growth activity; activin; inhibin activity; chemotactaxis; themokinetic activity; haemostasis; thrombolytic activity; receptor; ligand; anti-inflammatory; cadherin; tumour invasion suppressor;
(e.g. as vaccines) or suppressing activity, haematopoiesis regulating activity, tissue growth activity, activin/inhibin activity, activity, chemotactic/chemokinetic activity, haemostatic and thrombolytic activity, receptor/ilgand activity, anti-inflammatory activity, cadherin/tumour invasion suppressor activity, and tumour inhibition activity (no data is given in the specification to support these activities). The
                                                                                                                                     The present sequence represents a secreted protein. The polynucleotide and secreted protein are predicted to have biological activities which would make them suitable for treating, preventing or ameliorating medical conditions in humans and animals, although no supporting data is given. Suggested activities include nutritional activity, immune stimulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-MAR-1998;
21-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Secreted protein; nutritional activity; immune stimulating; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Secreted protein
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                                                                                                                                                                                                                                                                                                                       New polynucleotides encoding secreted human proteins - derived from human foetal brain, adult brain, foetal kidney, placenta or adult pineal gland cDNA libraries.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-OCT-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tumour inhibition;
                                                                                                                                                                                                                                                                             Claim 17; Page 73–74; 113pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1998-609890/51.
N-PSDB; AAV82780.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Racie LA,
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                                                                                                                                                                                                                                                                                                                                                                 AAU00512 standard; Protein; 298
                                                                                                           Doma in
                                                                                                                                                                                                                       Homo sapiens
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(TEXA-) TEXAS BIOTECHNOLOGY CORP.
                                     23-AUG-2000;
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                                     2000WO-US23158
                                                                                                           /note- ":
237 .254
                                                                                                                                                                             /note= "Possible signal peptide #1"
                                                                                                                                        /note=
                                                                                                                                                           /note= "Possible signal per ide #2"
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                                                                                                                                         "Possible mature JA:
                                                                                                 "Transmembrane domain"
                                                                                                                    "Possible mature JAM2 #2"
                                                                                                                                                                                                                                                                                                      protein (JAM2).
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Pred. No. 4.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The polynucleotide encoding the polypeptide is useful for recombinant production of JAM-2 protein, which in turn is useful for the production of antibodies. The antibodies may be used for probing cellular localisation and/or expression of JAM2 in tissues under normal and disease states, for immunoprecipitating JAM2 protein from cells and/or stroke tissues to determine whether it is modified by glycosylation and phosphorylation, and for determining JAM2 function. The antibodies and the protein of JAM2 with inflammatory cells or influences their aracellular migration, and is therefore useful for alleviating inflammatory diseases such as arthritis, asthma, rheumatoid arthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel nucleic acids encoding human junctional adhesion protein useful for producing antibodies that are suitable for therapeutic purposes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cunningham
immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma; inflammatlon; ischaemic shock; Alzheimer's disease; restenosis; AIDS; cognitive disorder; schizophrenia; prostate; obesity; osteoclast; thyrosteoporcsis; arthritis; testis; lung; thyroiditis; thyroid; digestion
                                                                      developmental abnormality; foetal deficiency; blood; allergy; renal;
                                                                                           diagnosis; tissue; cancer;
                                                                                                            Human;
                                                                                                                                                Human secreted
                                                                                                                                                                                    29-JAN-1999
                                                                                                                                                                                                                                                          AAW75220 standard; Protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence represents a human junctional adhesion molecule 2 (JAM2). polynucleotide encoding the polypeptide is useful for recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001-218425/22
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                                                                                                                                                                                                                                                                                                                                                                                                                        mqvddlnisgiiaavvvvalvisvcglgvcyaqrkgyf--ske-----tsfqksnssska
                                                                                     secreted protein; fusion protein; gene therapy; protein therapy; sis; tissue; cancer; tumour; neurodegenerative disorder; leukaemi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 46-47;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                298 AA;
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37.3%;
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                                                                                                                                              encoded by gene 25 clone HTEEB42
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                                                                                                                                                                                                                                                          298
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Pred. No. 4.3e-33;
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 digestion;
                                                                                           leukaemia;
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                                                                                                                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                                                                clone detailed in the descriptor line. The gene can be used to generate fusion proteins by linking to the gene to a human immunoglobulin Fc portion (e.g. AAV34277) for increasing the Lability of the fused protein as compared to the human protein only.

The invention relates to 28 novel genes and their fragments (nucleic acid sequences: AAV34286-V34325; amaino acid sequences AAV75196-W75235) which are useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. Also, pathological conditions can be diagnosed by determining the amount of the new polypucleotides. Specific uses are described for each of the 28 polynucleotides, based on which tissues they are most highly expressed in the polynucleotides, based on which tissues they are most highly expressed in the conditions can be diagnosed by determining the presence of mutations in the new polynucleotides. Specific uses are described for each of the 28 polynucleotides, based on which tissues they are most highly expressed in the conditions of the conditions are described for each of the 28 polynucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Isolated human poly:nucleotide(s) encoding secretory peptide(s) used to develop products for the diagnosis and treatment of e.g. inflammation, cancers, CNS disorders or immane system disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1998-520811/44.
N-PSDB; AAV34310.
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                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence represents a secreted human protein encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 168-169; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ferrie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-MAY-1997;
06-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-MAY-1997;
30-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-MAR-1997;
14-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HUMA-) HUMAN GENOME SCI INC.
              116
                                              52
                                                                             56
EVVALNDR-KEVDEITIELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWY
                                            tpkktvxsrlewkkl-grsvsfvyygqtlggdfknraemi-dfniriknvtrsdagkyrc
                                                                                                        marrsrhrl-----lllllrylvvalgyhkaygfsapkdqqvvtavxyqeailac-k 51
                                                                                                                                                                                                                                                                                    AAV34286 for described uses).
                                                                            THSQTSDPRIEWKKIQDGQTTYVYFDNKIQGDLAGRTDVIGKTSLRIWNVTRSDSAIYRC
                                                                                                                                            LSRRLRLYARLPHFFLLLLFRGCMI-----EAVNLKSSNRNPVVH--EFESVELSCII 55
                                                                                                                                                                        al Similarity
117; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Young
                                                                                                                                                                                                                                                      298 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fischer CL, Gentz KL,
Moore PA, Rosen CA,
                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Moore
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97US-0040710.
97US-0040762.
97US-00481100.
97US-0048189.
97US-0048357.
97US-0048970.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /label= unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'label= unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zeng Z;
                                                                                                                                                                                         30.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61;
                                                                                                                                                                          Score 498; DB 19;
Pred. No. 1.6e-32;
1; Mismatches 112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RĿ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ruben SM,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kyaw H;
                                                                                                                                                                                                        Length 298;
                                                                                                                                                                            Indels
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                                                                                                                   Query Match
Best Local
                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunoglobulin superfamily; Ig Sf; vascular adhesion molecule; inflammation; cancer; wound; angiogenesis; human; confluency regulated adhesion molecule 2; CRAM-2; JAM-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAB27273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB27273 standard; Protein; 298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human confluency regulated adhesion molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 110
                                                                                                                                                                                                The present sequence is the human confluency regulated adhesion molecule 2 (CRAM-2, also known as JAM-3). CRAM-2 is one of the vascular adhesion proteins of the immunoglobulin superfamily (Ig Sf). The CRAM-2 protein and coding sequence can be used in the treatment of cancer, inflammation, to modulate cell interactions and angiogenesis, and in the modulation of wound healing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200053749-A2
                                                                                                                                                                                                                                                                                                                       Isolated human Confluency Regulated Adhesion Molecule 1 or 2 (CRAM-1 or CRAM-2) polypeptide, useful for treatment of tumors, inflammation reactions and modulating vascular permeability .
                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-MAR-2000; 2000WO-EP02219
                                                                                                                                                                                                                                                                                                                                                                                                        Imhof BA, Aurrand-Lions M;
                                                                                                                                                                                                                                                                                                                                                                                                                                  (RMFD-) RMF DICTAGENE SA.
                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                               aim 1; Fig 3; 59pp; English
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 61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEVYDLNIAGIIGGVLVVLIVLAVITMGICCAYRRGCFISSKQDGESYKSPGKHDGVNYI 294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            \verb|evsapsegggnleedtvtlevlvapavpscevpssalsgtvvelrcqdkegnpapeytwf|
                                                      2
                                                                            ARLPHFFLLLLFRGCMIEAVNLKSSN------RNPVVHEFESVELSCIITHSQTSDP 63
                         RIEWKKIQDGQTTVVYFDNKIQGDLAGRTDVFGKTSLRIWNVTRSDSAIYRCEVVALNDR 123
                                                     117;
rlewkkvgqg-vslvyyqqalqgdfkdraemi-dfniriknvtrsdageyrcevsapteq 118
                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                           298 AA;
                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99EP-0200746
                                                                                                          30.0%; Score 490.5; DB 21; 37.6%; Pred. No. 6.3e-32; tive 49; Mismatches 118;
                                                                                                                 Indels
                                                                                                                                       Length 298;
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13

ARLPHEFLLLLERGCMIEAVNLKSSN-----RNP. VHYFESVELSCIITHSQTSDP 63

Indels

27;

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Query Match Best Local Similarity

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64 RIBWKKIQDGQTTYVYFDNKTQGDLAGRTDVFGKTSI.5 "...NVTRSDSAIYRCEVVALNDR 123

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunoglobulin superfamily; Ig Sf; vasculo: Jdhesion molecule; inflammation; cancer; wound; angiogenesis; Juse; confluency regulated adhesion molecule 2; CkAN-2; JAM-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Murine confluency regulated adhesion molecule 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB27275 standard; Protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     119
                                                                                         The present sequence is the murine confluence in caulated adhesion molecule 2 (CRAM-2, also known as JAM-3). CRAM-2 is one of the vascular adhesion proteins of the immunoglobulin superfamily (19 Sf). The CRAM-2 protein and coding sequence can be used in the treatment of cancer, inflammation, to modulate cell-cell interactions and angionesis, and in the modulation of wound healing.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated human Confluency Regulated Adhess. Molecule 1 or 2 (CRAM-2) polypeptide, useful for treatment c. tumors, inflammation reactions and modulating vascular permeability
                                                                                                                                                                                                                                                                                                                                                            11-MAR-1999;
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                                                                                                                                                                                        Example; Fig 5; 59pp; English.
                                                                                                                                                                                                                                                                           WPI; 2000-587436/55
                                                                                                                                                                                                                                                                                                      Imhof BA,
                                                                                                                                                                                                                                                                                                                                (RMFD-) RMF DICTAGENE SA
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                                                                  Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gqnlqedkvmlevlvuFavpacevptsvmtgsvvelrcijdkcynpupoviwfkdg----t 174
                                                                      298 AA:
130.0%; Score 490.5; ... 21; ilarity 37.6%; Pred. No. 6.3e 42; Conservitive 49; Mismatches 118;
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                                 21; Length 298;
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             This invention describes a novel composition containing (apart from a carrier or excipient), a novel PRO245 polypeptide (I), its agonist or antagonist, or their fragments, for modulating: (i) infiltration of inflammatory cells into tissue; (ii) an immune response; or (iii) T co
                                                                                                                                                                                                                     Fong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Inflammatory cell infiltration; immune response; T cell proliferation; anti-inflammatory; anti-autoimmune; anti-diabetic; spondyloarthropathy; T cell-mediated disease; spondyloarthropathy; sclerosis; renal disease; inflammatory myopathy; hemolytic anemia; thrombocytopenia; thyroiditis; inflammatory myopathy; hemolytic anemia; thrombocytopenia; thyroiditis; diabetes mellitus; demyelinating polyneuropathy; currhosis; enteropathy; multiple sclerosis; polyneuropathy; hepatitis; cirrhosis; enteropathy; hepatitis; inflammatory bowel disease; whipple; disease; sclerosing cholangitis; inflammatory bowel disease; whipple; disease;
                                                                                           Example 1; Fig 2;
                                                                                                                          antagonist
                                                                                                                                       Composition containing novel polypeptide PRO245, its
                                                                                                                                                                                                                                                                                                  21-NOV-1997;
24-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9914241-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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AAY08060 standard; Protein; 312 AA.
                                                                                                                                                                                                                                                   (GETH )
                                                                                                                                                                                                                                                                                                                                                 28-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                               17-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     idiopathic pulmonary fibrosis; graft rejection; PRO245; human.
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                                                                                                                                                                        1999-229499/19
DB; AAX37664.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease; dermatitis; psoriasis; asthma; allergic rhinitis; tumor hypersensitivity; urticaria; eosinophilic pneumonia; transplant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ndfrhtksfii 298
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              lnisgiiatvvvvafvisvcglgtcyaqrkgyf--sketsfqkgsp-----askvttmge 287
                                                                                                                                                                                                                                                    GENENTECH INC
                                                                                                                                                                                                                     Goddard A,
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97US-0059263.
97US-0063550.
97US-0065186.
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97US-0066770.
98US-0088026.
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                                                                                        177pp; English
 composition increases
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(iii) T cell
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Best Local S
Matches 110
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            bullous skin diseases, erythema multiforme contact dermutitis, psoriasis, asthma, allergic rhinitis, atopic dermatitis, food hypersensitivity, urticaria, eosinophilic pneumonia, idiopathic pulmonary fibrosis, hypersensitivity pneumonitis, and transplantation associated diseases (graft rejection, and graft versus-host-disease). (1), its (ant)agonists or fragment can also be used as an adjuvant in treatment of tumors. Antibodies against (1) can also be used for diagnosing such diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hemoglobinuria), autoimmune thrombocytopenia (idiopathic thrombocytopenic purpura immune-mediated thrombocytopenia), thyroiditis (Grave's disease, Hashimoto's thyroiditis, juvenile lymphocytic thyroiditis, atrophic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              arthritis, spondyloarthropathies, systemic sclerosis (scleroderma), idiopathic inflammatory myopathies (dermatomyositis, polymyositis), Sjogren's syndrome, systemic vasculitis, sarcoidosis, autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and their fragments, are used particularly T cell-mediated c
                                                                                                                                                   AAY23324 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (i)-(iii). The products of the invention have anti-inflammatory, and its (anti-agonists
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hemolytic anemia (immune pancytopenia, paroxysmal nocturnal
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                                                                                                                                                                                                                                                                                                                                                                       292 NYI 294
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nes 110; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence represents the human PRO245 protein described in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEVYDLNIAGIIGGVLVVLIVLAVITMGICCAYRRGCF - -- ISSKQDGESYKSPGKHDGV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              marrsrhrl-----lllllrylvvalgyhkayolsapkdqqvvtaveyqeailac-k
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          kdgirllenprlgsqstnssytmntktgtlqfntvskldtgeyscearnsvgyrrcpgkr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EVVALNDR-KEVDEITIELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWY 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     THSQTSDPRIEWKKIQDGQTTYVYFDNKIQGDLAGKTDVFGKTSLRIWNVTRSDSAIYRC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               312 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ments, are used to treat immune-related diseases, cell-mediated diseases. The diseases treated include
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29.3%; Score 478; DB 20; 36.3%; Pred. No. 6.9e-31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            62;
                                                                                                                                                      312
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A33 related antigen PRO245

02-SEP-1999

(first entry)

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

OM protein - protein search, using sw model

Run on:

August 6, 2001, 09:31:29; Search time 25.41 Seconds (without alignments) 929.324 Million cell updates/sec

Title: Perfect score: US-09-524-531A-13 1633

Sequence:

1 MALSRRLRLYARLPHFFL..........VNYIRTSEEGDFRHKSSFVI 310

Scoring table:

hed:

BLOSUM62 Gapop 10.0 , Gapext 0.5

number of hits satisfying chosen parameters:

219241

219241 seqs, 76174552 residues

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database : PIR_68:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	80	7	į	5	4		20)_	No.	Result
151.5	152	153	153.5	154.5	156	156	156	157	159.5	160	162	162	163	164	164.5	164.5	166	167.5	167.5	169	169	169.5	169.5	172.5	177	178	180	181	Score	
9.3	9.3	9.4	9.4	9.5	9.6	9.6	9.6	9.6	9.8	9.8	9.9	9.9	10.0	10.0	10.1	10.1	10.2	10.3	10.3	10.3	10.3	10.4	10.4	10.6	10.8	10.9	11.0	11.1	Match	Query
811	538	1273	1033	272	4162	1323	344	1091	1277	875	5198	5175	761	7962	853	352	483	6642	3707	1092	725	1115	725	858	333	1051	1088	725	Length	
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A41054	JC2457	T42405	S19247	I48268	T42633	PN0568	156551	IJCHNL	T30532	T33434	T43290	T20992	IJHUNG	I38346	IJBONC	T33433	T17346	T29757	S18252	JN0635	JE0100	IJMSNL	IJMSNG	IJRTNC	A31923	A39712	IJXLNL	JE0099	ij	
fasciclin II, tran	vascular cell adhe	sax-3 protein - Ca	cell adhesion prot	biliary glycoprote	connectin/titin -	connectin 3B - chi			neural cell adhesi	hypothetical prote	hemicentin precurs	hypothetical prote	neural cell adhesi	elastic titin - hu			hypothetical prote	UNC-89 -	heparan sulfate pr	neural cell adhesi	l cell	cell	cell	(2		-like	neural cell adhesi	cell adhe	Description	

RESULT 2
LOXLINL
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unc-5 protein, lor axonal glycoprotei transient axonal c		•				
procein-c axonal gl	24556	د	1040	۰	148	
unc-5 pro	A34695	N	1040	9.1	148	
process.	B44294	_	947	9.1	148	
110+041-4	JC4593	N	1070	9.1	148.5	
protein-t	I48697	_	881	9.1	148.5	
protein-tyrosine k	148696	_	871	9.1	148.5	
dutt1 protein -	T30805	N	1612	9.1	149	
perlecan precurson	A38096	N	4391	9.2	149.5	
sialoadhe	S50065	N	1694	9.2	150	
hypothetical prote	T08678	N	584	9.2	150	
hypothetical	T29549	N	423	9.2	150	
protein-tyrosine-	A56178	N	1912	9.2	151	
protein-tyrosine-p	C54689	N	1894	9.2	151	
glial cell membrar	A58532	N	1091	9.2	151	
tumor suppressor	151669	N	1427	9. 3	151.5	
fasciclin II PI-li	B41054	N	873	9.3	151.5	

ALIGNMENTS

Db 264 QSEMTIHHVEKDDEAEYSCIANNQAGEA 291	Qy 201 TGTLVENAVHKDDSGQYYCIASNDAGAA 228	Db 213 ARQLRVNATAKMAESVVLSC-DADGFPDPEISWLKKGEPIEDGEEKISFNED 263	162RFVVLANNYLQIRGIKKTDEGTYRCEGRILARGEINYKDIQVIVNVPPTIQ	Qy 32 VNLKSNRNPVHEFSSVELSCIITHSQTSDPR-IEWK-KIQDGTTYVYEDKI 84	Query Match 11.1%; Score 181; DB 2; Length 725; Best Local Similarity 28.8%; Pred. No. 6.3e-07; Matches 60; Conservative 33; Mismatches 81; Indels 34; Gaps 11;	RESULT 1 JE0099 neural cell adhesion molecule 1 - African clawed frog N;Alternate names: N-CAM 1 C;Species: Xenopus laevis (African clawed frog) C;Date: 19-May-1998 #sequence_revision 29-May-1998 #text_change 21-Jul-2000 C;Accession: JE0099 R;Kudo, M.; Takayama, E.; Tadakuma, T.; Shiokawa, K. Biochem. Biophys. Res. Commun. 245, 127-132, 1998 A;Title: Molecular cloning of ssd-form neural cell adhesion molecules (N-CAMs) as the A;Accession: JE0099; MUID:98204770 A;Accession: JE0099 A;Molecule type: mRNA A;Residues: 1-725 <kud> A;Cross-references: DDBJ:AB008162; NID:g3116226; PIDN:BAA25931.1; PID:g3116227 A;Experimental source: heart C;Comment: This protein mediates and regulates various cell-cell interactions through C;Superfamily: neural cell adhesion molecule; fibronectin type III repeat homology; 1 F;512-589/Domain: fibronectin type III repeat homology; 1</kud>
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A;Molecule type: mRNA
A;Residues: 1-1051 <CHO>
A;Residues: 1-1051 <CHO>
A;Cross-references: GB:M63437; NID:g212235; PIDN:AAA48933.1; PID:g212236
C;Superfamily: unassigned Ser/Thr or Tyr-specific protein kinases; protein C;Keywords: ATP
E;775-1046/Domain: protein kinase homology <KIN>
F;7783-791/Rayion: protein kinase ATP-binding motif
                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT
A39712
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: mRNA
A; Residues: 1-1088 < KRI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic Acids Res. 17, 10341-10333, 1707
A;Title: Primary structure and developmental expression of a large cytoplasmic
A;Reference number: S09600; MUID:90098871
                                                                                                                                                                                                        A; Reference number: A; Accession: A39712
                                                                                                                                                                                                                                 A; Title: Characterization of A; Reference number: A39712; N
                                                                                                                                                                                                                                                                                                                            kinase-like protein klg precursor - chicken
C;Species: Gallus (chicken)
C;Date: 08-Nov-1991 #sequence_revision 08-Nov-1991 #text_change 24-Sep-1999
C;Accession: A39712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Gene: NCAM
                                                                                                                                                                                 A;Status: preliminary
                                                                                                                                                                                                                                                                                 Proc. Natl. Acad.
                                                                                                                                                                                                                                                                                                     R;Chou, Y.H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   F;724-1088/Domain:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;Genetics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        381/Domain: immunoglobulin homology <IMM4>
4.5-475/Domain: immunoglobulin homology <IMM5>
512-589/Domain: ifibronectin type III repeat homology <FN3A>
;618-679/Domain: fibronectin type III repeat homology <FN3B>
;706-723/Domain: transmembrane #status predicted <FNM>
;723-723/Domain: intracellular #status predicted <INT>
;74-1088/Domain: intracellular #status predicted <INT>
;41-93,136-186,232-282,333-379,420-473/Disulfide bonds: #status predicted
;41-93,136-186,232-282,333-379,420-473/Disulfide bonds: #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1-19/Domain: signal sequence #status predicted <SIG>
20-1088/Product: neural cell adhesion molecule, long domain for 20-803,1050-1088/Product: neural cell adhesion molecule, short
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Superfamily: neural cell adhesion molecule; fibronectin type III repeat homology; is Keywords: alternative splicing; brain; cell adhesion; duplication; heparin binding;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     188/Domain:
153/Region:
158-162/Region:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             143 V--CRIPAAVPVGKTATLQCQESEGYPRPHYSWYRNDVPLPTDSRANPRFQNSSFHVNSE 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           284/Domain:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05/Domain: extracellular #status predicted <EXT>5/Domain: immunoglobulin homology <IMM1>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                85
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           QGDLAGRTDVFGKTSLRIWNVTRSDSAIYRCE--VVALNDRKEVDEITIELIVQVKPVTP 142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VNLKIYQKLTFKNAPTPQEFKEGEDAVIICDVSSSIPSIITWRHKGKD-----VIFKKDV 161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  QSEMTIHHVEKDDEAEYSCIANNQAGEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TGTLVFNAVHKDDSGQYYCIASNDAGAA 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ARQLRVNATANMAESVVLSC-DADGFPDPEISWLKKGEPIEDGE-----EKISF--NED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -----RFVVLANNYLQIRGIKKTDEGTYRCEGRILA----RGEINYKDIQVIVNVPPTIQ 212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              60;
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heparin binding #status predicted
heparin binding #status predicted
heparin binding #status predicted
immunoglobulin homology <IMM3>
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f a member of the
MUID:91271300
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Pred. No. 1.2e
33; Mismatches
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                                                                                                                                                                                                                                                        1991
immunoglobulin
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A; Accession: A31923
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Species: Drosophila melanogaster
C;Date: 18-Oct-1989 #sequence_revision 18-Oct-1989 #text_change
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neural cell adhesion molecule short domain N; Alternate names: NCAM-140 C; Species: Rattus norvegicus (Norway rat) C; Date: 30-Sep-1991 #sequence_revision 30-5

30-Sep-1991

#text_change

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JAM1_MOUSE
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Best Local
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EMBL; U89915; AAC32982.1; -. MGD; MGI:1321398; Jcam. InterF. IPR003006; -.
                                                                                                            This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE-98327120;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus
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01-0CT-2000 (Rel. 40, Last sequence up
01-0CT-2000 (Rel. 40, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JAM1_MOUSE 088792;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=10090;
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                                                                                                                                                                                                                                                                                                                                 ACTIVATION.
SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN (POTENTIAL).
TISSUE SPECIFICITY: LOCALIZED AT TIGHT JUNCTIONS OF BOTH
EPITHELIAL AND ENDOTHELIAL CELLS.
SIMILARITY: CONTAINS 2 IMMUNOGLOBULIN-LIKE V-TYPE DOMAINS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RTSEEGDFRHKSSFVI 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EVSAPSEQGQNLEEDTVTLEVLVAPAVPSCEVPSSALSGTVVELRCQDKEGNPAPEYTWF
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Chordata;
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MOLECULE PRECURSOR (JAM).
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Pred. No. 4.1e-37;
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Best Local S
Matches 102
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Tight junction;
Signal.
SIGNAL 1
**Rornecki E.;

**Molecular cloning and sequencing of the cDNA of F11 receptor, novel Ig superfamily member from human platelets.";

Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases.
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DISULFID
CARBOHYD
CARBOHYD
                                                                                                                                                                                                                                                                                                                        JAM1_HUMAN STANDARD;
Q9Y624;
01-OCT-2000 (Rel. 40, Created)
01-OCT-2000 (Rel. 40, Last seq
01-OCT-2000 (Rel. 40, Last ann
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DOMAIN
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                            01-OCT-2000 (Rel. 40, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
JUNCTIONAL ADHESION MOLECULE PRECURSOR (JAM)
                                                                                                                                                                         MEDLINE=99323940; PubMed=10395639; Ozaki H., Ishii K., Horiuchi H., A
                                                                                                                                                                                                                                                                                    JCAM OR JAM1.
                                                                                                                                                                                                                                                                                                MOLECULE 1) (PAM-1) (PLATELET F11 RECEPTOR).
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                                                                                SEQUENCE FROM
                                                                                                          J. Immunol. 163:553-557(1999).
                                                                                                                                  redistribution of junctional adhesion
                                                                                                                                                                                                      SEQUENCE FROM
                                                                                                                                                                                                                                                                      Homo sapiens (Human).
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                                                                                                                                   "Cutting edge: combined treatment of TNF-alpha and IFN-gamma causes redistribution of junctional adhesion molecule in human endothelial
                                                                                                                                                              Iwamatsu A., Kita T.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 247
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LLFLFTSMILGSLVQGKGSVYTAQSDVQVPENESIKLTC--TYSGFSSPRVEWKFVQGST 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LLLLFRGCMIEAV-----NLKSSNRNPVVHEFESVELSCIITHSQTSDPRIEWKKIQDGQ 74
                                                                                                                                                                                                                                                                                                                                                                                                                                               LILLGLLIFGVWFAYSRGYFETTKKG----TAPGKKVIYSQPSTRSEGEFKQTSSFLV
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102; Conserv
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IG-LIKE V-TYPE DOMAIN.
IG-LIKE V-TYPE DOMAIN.
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POTENTIAL.
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N-LINKED (GLCNAC. . .) (POTENTIAL).
391F3E48FF3B97EC CRC64;
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; Mismatches
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                           Minimum DB seq length: 0
Maximum DB seq length: 200000000
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Perfect score:
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Gapop 10.0 , Gapext 0.5
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1633
1 MALSRRLRLRLYARLPHFFL.....VNYIRTSEEGDFRHKSSFVI 310
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Database : SPTREMBL_16:*
1: sp_archea: sp_plant:*
sp_rodent:*
sp_unclassified:*
sp_vertebrate:*
sp_virus:* sp_archea:*
sp_bacteria:*
sp_fungi:* sp_organelle:*
sp_phage:* sp_mhc:* sp_human:* sp_mammal:* sp_invertebrate:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

19	18	17	16	15	14	13	12	11.	10	9	ω,	7	6	. U	4	w	2	_	No.	•
174.5	177	181	182.5	197	199	208	209.5	212	213	214	219	240.5	257	321	340	450.5	497.5	1615	Score	
10.7	10.8	11.1	11.2	12.1	12.2	12.7	12.8	13.0	13.0	13.1	13.4	14.7	15.7	19.7	20.8	27.6	30.5	98.9	Query Match Length	a s
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Q9tu80 canis famil	Q9v3a5 drosophila	073633 xenopus lae	Q90490 brachydanio	095532 homo sapien		Q9ygv5 gallus gall	Q9yghl gallus gall	Q9z109 mus musculu	Q9nx42 homo sapien	Q9pwr4 gallus gall	095791 homo sapien	Q9jka5 mus musculu	Q91664 xenopus lae	09jkd5 rattus norv	Q9y5b2 homo sapien	Q9jhy1 rattus norv	Bug	Q9epk4 mus musculu	Description	

443 443 443	39 40 11	365	3333 4331	27 28 29	20 21 22 23 24 26
158.5 157 157 157 157	159.5 158.5 158.5	160	163 162 161	164.5 164.5 164 163	172.5 169 168.5 168.5 167.5 166
9.667	9.7	9999 8888	10.0 9.9 9.8	10.1	10.6 10.3 10.3 10.3 10.3
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09eqs8 mus musculu 057596 gallus gall 09dg15 gallus gall 093242 gallus gall	098902 fugu rubrip 09tu79 sus scrofa 018466 hirudo medi 09eqs9 mus musculu	076698 caenorhabdi Q9u4d1 caenorhabdi Q9u4d1 caenorhabdi Q9v4y0 drosophila	homo caen dros		Q9ukv4 homo sapien 073634 xenopus lae 09vwz7 drosophila Q61349 mus musculu 001761 caenorhabdi 09uf14 homo sapien 092626 homo sapien

ALIGNMENTS

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181 PTDSRANPREQNSSEHVNSETGTLVENAVHKDDSGQYYCIASNDAGAARCEGQDMEVYDL 240	121 NDRKEVDEITIELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWYRNDVPL 180 	61 SDPRIEWKKIQDGQTTYVYFDNKIQGDLAGRTDVFGKTSLRIWNVTRSDSAIYRCEVVAL 120 	1 MALSRRLRLRLYARLPHFFLLLLFRGCMIEAVNLKSSNRNPVVHEFESVELSCIITHSQT 60	Query Match 98.9%; Score 1615; DB 11; Length 310; Best Local Similarity 99.4%; Pred. No. 2.2e-153; Matches 308; Conservative 0; Mismatches 2; Indels 0; Gaps 0;	EMBL; AJ300304; CAC20704.1; SEQUENCE 310 AA; 34837 MW; 4B92BCB51D0A4B0A CRC64;	Curr. Top. Microbiol. Immunol. 251:91-98(2000).		"Cloning of JAM-2 and JAM-3: an Emerging Junctional Adhesion Molecular	Aurrand-Lions M.A., Duncan I., Du Pasquier I., Tmhof B.A.;	SEQUENCE FROM N.A.	[1]			Fukbarvota: Metazoa: Chordata: Craniata: Vertebrata: Euteleostomi:	Miss Buscolline (Moliso)	JUNCTIONAL ADHESION MOLECULE-Z, JAM-Z. JAM-Z.	01-MAR-2001 (TrEMBLrel, 16, Last annotation update)	16,	2001 (TrEMBLrel.		N4 OGEDKA PRELIMINARY: PRT: 310 AA	IT 1

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Q9JI59;
01-OCT-2000
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J. Biol. Chem. 275:19139-19145(2000).
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Rukaryota; Metazoa; Chordata;
ammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
VASCULAR ENDOTHELIAL JUNCTION-ASSOCIATED MOLECULE (JUNCTIONAL ADHESION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR003598; -. Pfam; PF00047; 1g; 2. SMART; SM00408; IGc2; 1
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Aurrand-Lions M.A., Duncan L., I
"Cloning of JAM-2 and JAM-3: an
Family?";
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MEDLINE-20317114; PubMed-10779521;
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               235
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                                                                                                         SLLGNPKGGTHNNSSYTMNTKSGILQFNMISKMDSGEYYCEARNSVGHRRCPGKRMQVDV 234
                                                                                                                                                         DSRANPR----FQNSSFHVNSETGTLVFNAVHKDDSGQYYCIASNDAGAARCEGQDMEVYD
                                                                                                                                                                                                          GQNLQEDKVMLEVLVAPAVPACEVPTSVMTGSVVELRCQDKEGNPAPEYIWFKDG----T
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CE 298 AA;
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33047 MW;
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Pred. No. 1.5e-41;
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Best Local
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Submitted (JUN-2000) to the E
EMBL; AF276998; AAF78250.1; -
InterPro; IPR003006; -
InterPro; IPR003596; -
Pfam; PF00047; 19; 2
SMART; SM00406; IGv; 1.
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Q9JHY1;
Q1-OCT-2000 (TrEMBLrel. 15, Created)
Q1-OCT-2000 (TrEMBLrel. 15, Last sequence update)
Q1-MAR-2001 (TrEMBLrel. 16, Last annotation update)
JUNCTIONAL ADHESION MOLECULE JAM.
                                                                                                     01-NOV-1999 (TrEMBLrel. 12, 01-NOV-1999 (TrEMBLrel. 12, 01-MAR-2001 (TrEMBLrel. 16, JUNCTION ADHESION MOLECULE.
                                                                                                                                         Q9Y5B2 PRELIMINARY;
Q9Y5B2;
Q1-NOV-1999 (TrEMBLrel.
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SEQUENCE FROM N.A.
Liu Y., Nusrat A., Schnell
Foley C., Parkos C.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                    NCBI_TaxID=9606;
                                                                Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                         Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
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Local Similarity 38.0%;
nes 114; Conservative 5
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                                                                                                                                                                                                                                     LILLGLLIFGIWFAYSRGYFERTKKG----TAPGKK--VIYSQPSARSEGEFKQTSSFLV
                                                                                                                                                                                                                                                             LIVLAVITMGICCAYRRGCFISSKQDGESYKSPGKHDGVNYIRTS--EEGDFRHKSSFVI 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LLLLERGCMI-EAVNLKSSNRNP----VVHEFESVELSCIITHSQTSDPRIEWKKIQDGQ
                                                                                                                                                                                                                                                                                       {\tt SYTIDPKSGDLVFDPVSAFDSGEYYCEAQNGYGTAMRSEAVRMEAVELNVGGIVAAVLVT}
                                                                                                                                                                                                                                                                                                     SFHVNSETGTLVFNAVHKDDSGQYYCIASNDAGAA-RCEGQDMEVYDLNIAGIIGGVLVV
                                                                                                                                                                                                                                                                                                                                          VLVPPSKPTVSIPSSVTIGNRAVLTCSEHDGSPPSEYSWFKDGVPMLTADAKKTRAFINS
                                                                                                                                                                                                                                                                                                                                                         VQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWYRNDVPLPT-DSRANPRFQNS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GDFRHKSSFVI 310
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                300 AA;
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                                                                Chordata;
Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                32369 MW;
              F.J.,
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                                                                                                                 Last sequence up
                                                                                                                                            Created)
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                                                                Craniata; Vertebrata; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Craniata; Vertebrata; Sciurognathi; Muridae;
                                                                                                                                                                       PRT;
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                Walsh
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              s.,
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                                                                                                                  on update)
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                Reaves
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            databases
              T.A.,
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                                                                  Euteleostomi;
; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Murinae;
                Pochet M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                        246
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August 6, 2001, 09:31:09; Search time 19.31 Seconds (without alignments) 330.554 Million cell updates/sec

Sequence: Title: Perfect score:

US-09-524-531A-13
1633
1 MALSRRLRLRLYARLPHFFL.....VNYIRTSEEGDFRHKSSFVI 310

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

ched: 197339 seqs, 20590346 residues

number of hits satisfying chosen parameters:

197339

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

Issued_Patents_AA:*

1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	6	5		3	2		Result
141	141	144.5	144.5	144.5	144.5	144.5	146.5	146.5	146.5	147	147.5	148	151	151.5	156.5	156.5	160	160	166	171.5	172.5	172.5	173.5	242.5	403	415	Score
8.6	8.6	8.8	8.8	8.8	8.8		9.0	9.0	9.0	9.0	•	9.1	9.2	9.3	9.6	9.6	9.8	9.8	10.2	10.5	10.6	10.6	10.6	14.8	24.7	25.4	Query Match Length
642	338	868	868	734	734	869	868	860	478	95	501	607	1091	365	365	365	869	869	1101	390	365	365	365	319	299	299	
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US-08-217-299-1	US-08-414-657D-60	US-08-644-271-1	US-08-374-834-1	US-08-987-867A-17	US-08-389-459A-17	US-08-602-725-36	PCT-US95-08493-21	PCT-US95-08493-19	PCT-US95-08493-15	US-08-928-383B-18	US-08-408-095-31	US-08-752-307B-12	US-08-986-485-5	US-08-928-383B-26	US-08-928-383B-24	US-08-928-383B-23	US-08-644-271-29	US-08-374-834-16	US-08-986-485-2	US-08-979-424-1	US-09-272-496-2	US-08-979-424-3	US-08-928-383B-2	US-08-597-495B-22	US-09-188-930-189	US-09-188-930-331	ID
1,	60,	1,	Sequence 1, Appli	17,	17,	36,	21,	19,	15,	18,	31,	12,		26,	24,	23,	29,	16,	'n	1,	Sequence 2, Appli	Sequence 3, Appli	Sequence 2, Appli	22,	•	Sequence 331, App	Description

RESULT

2

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135.5	138.5	138.5	138.5	138.5	138.5	138.5	138.5	138.5	138.5	139	139	139	139.5	139.5	139.5	141	141
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662	325	325	315	308	304	287	287	252	252	338	338	310	1447	1447	612	1501	1501
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-261	•	US-08-414-657D-2	US-08-414-657D-47	US-08-414-657D-46	US-08-414-657D-44	US-08-414-657D-49	US-08-414-657D-48	US-08-414-657D-57	US-08-414-657D-56	US-08-414-657D-43	US-08-414-657D-42	US-08-414-657D-45	PCT-US94-05277-2	US-09-041-886-25	US-08-752-307B-11	US-08-716-679-3	US-08-447-464-3
7, 7	4.	2,	Sequence 47, App.	•	Sequence 44, App.	Sequence 49, Appl	•	•	Sequence 56, App	•	-	Sequence 45, App.	Sequence 2, Appli	Sequence 25, App.	Sequence 11, App.	Sequence 3, Appl.	Sequence 3, Appl.

ALIGNMENTS

6;

Diskette, 3.5 inch,

360 kb storage

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                                                                                                                                                                                                                                                                                                                                    Sequence 22,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Watson, James D.
APPLICANT: Strachan, Lorna
APPLICANT: Sleeman, Matthew
APPLICANT: Onrust, Rene
APPLICANT: Murison, James Greg
APPLICANT: Murison, James Greg
TITLE OF INVENTION: Compositions Isolated From Skin Cells
TITLE OF INVENTION: and Methods For Their Use
FILE REFERENCE: 11000.1011c1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 189 LENGTH: 299
                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/188,930A CURRENT FILING DATE: 1998-11-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                  tent No.
COMPU
                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                           APPLICANT: Simpson, Richard J.; Nice, Edouard; Moritz, R. L.; APPLICANT: Catimel, B.; Ji, Hong; Burgess, Anthony W.; APPLICANT: Heath, Joan K.; White, Sara J.; Johnstone, Cameron
                                                                                                                                                                                                                                                        APPLICANT: Old, Lloyd J.; APPLICANT: Simpson, Richa:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AME/KEY: UNSURE
                                                                                                                                                                          TITLE OF INVENTION: Colon Cell Associated
                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ME/KEY: UNSURE
CATION: (289)...(289)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                     251 GILVFGIWFAYSRGHFDRTKKGTSSKK-----VIYSQPSARSEXEFKQTSSFLV 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                257
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                                  STATE: N
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                                                       STATE: New York City
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 33.1 nes 98; Conservative
                                                                                                STREET:
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                                                                                                                   ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NSETGTLVFNAVHKDDSGQYYCIASNDAGAARCEGQ-DMEVYDLNIAGIIGGVLVVLIVL 256
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILAILLCSLALGSVTVHSSEPEVRIPENNPVKLSC--AYSGFSSPRVEWKFDQGDTTRL 72
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             189, Application US/09188930A
3. 6150502
                    1002

    Application US/08597495B
5712369

READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (247)...(247)
                                                                                            3: Felfe & Lynch
805 Third Avenue
                                       USA
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Pred. No. 1.2e-32;
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                                                                                                                                                                                                                                                                          Sydney; Ritter,
                                                                                                                                                                            Molecules, Protein And Peptides
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                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: ROBERT W.
APPLICANT: and Marsh
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                              Sequence 2, Application US/08928383B Patent No. 6210921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 04-Aug-1995
ATTORNEY/AGENT INFORMATION:
NAME: Hanson, No. 571236
                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/511,876
FILING DATE: 04-Aug-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
COMPUTER READABLE FORM:
                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                         294 EREEEDDYRQE 304
                                                                                                                                                                                                                                                                                                                                                                                            295 RTSEEGDFRHK 305
                                                                                                                                                                                                                                                                                                                                                                                                                                239
                                                                                                                                                                                                                                                                                                                                                                                                                                                             239 DLNIAGIIGGYLYVLIVLAVITMGICCAYRRGCFISSKQDG----ESYKSPGKHDGVNYI 294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    194
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        134 RLLVLVPPSKPECGIEGETIIGNNIQLTCQSKEGSPTPQYSWKRYNILNQEQPLAQPASG 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/597,495B FILING DATE: 02-Feb-1996 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19
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nes 80; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 30, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE:
COMPUTER: II
                                 COUNTRY:
                                                                      CITY:
                                                                                     STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           78 NYIHGELYKNRVSISNNAEQSD----ASITIDQLTMADNGTYECSVSLMSDLEGNTKSRV 133
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                                                   STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29 IEAVNLKSSNRNPVVHEFESVELSCIITHSQTS--DPRIEWKKIQDGQTTYVY---FDNK 83
                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               POPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                              ----VGIAVGVVAALIIIGII-IYCCCCRGKDDNTEDKEDARPNREAYEEPPEQLRELSR 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWYRNDV-----PLPTDSRA 186
                    02109
                                                                    Boston
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                                                 Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             319 amino acids
                                   USA
                                                                                      28 State Street
                                                                                                                                                                          and Marshall S. Hor
                                                                                                                                                                                                           Robert W. Finberg, Jeffrey M. Bergelson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (212)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wordperfect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IBM PS/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (212) 688-9200
                                                                                                       LAHIVE & COCKFIELD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    838-3884
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14.8%;
25.7%;
                                                                                                                                                         Receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -VSLKNISTDTSGYYICTSSNEEGTQFCNITVAVRSPSMNVALY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30,946
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       49;
                                                                                                                                                                                              Horwitz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LUD 5316.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 242.5; DB 1
Pred. No. 1.9e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                             6210921el Coxsackievirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 319;
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                                                                                                                                                                             and
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